

SEQUENCE LISTING

<110> Phagetech, Inc. PELLETIER, Jerry GROS, Phillippe DUBOW, Michael

- <120> DNA SEQUENCES FROM STAPHYLOCOCCUS AUREUS BACTERIOPHAGE 44 AHJD THAT ENCODE ANTI-MICROBIAL POLYPEPTIDES
- <130> 073406-0302
- <140> 09/727,892
- <141> 2000-12-01
- <160> 159
- <170> PatentIn version 3.0
- <210> 1
- <211> 16668
- <212> DNA
- <213> Staphylococcus aureus Bacteriophage 44 AHJD
- <400> 1

tccatttctt tactaaactt aaaaatgctg tgcaacaact taaccaactt atctaaccta 60 ttacatattc atcaaataca aaatttatgt atctattgac ttttattcaa aattatgatt 120 tcaacatata ataaaattaa tttacttatt taaatattct atgatataat tagttataaa 180 atatttggag gtgtataaat gacagaattt gatgaaatcg taaaaccaga cgacaaagaa 240 qaaacttcag aatcaactga agaaaattta gaatcaactg aagaaacttc agaatcaact 300 qaaqaatcaa ctgaaqaatc aactgaaqaa tcaactgaag ataaaacagt agaaacaatc 360 qaaqaaqaaa atgaaaacaa attagaacct actacaacag atgaagatag ttcgaaattt 420 gaccctgttg tattagaaca acgtattgct tcattagaac aacaagtgac tactttttta 480 540 tcttcacaaa tgcaacaacc acaacaagta caacaaacac aatcagatgt aacagaatca aacaaagaag ataacgacta ttcagatgaa gaactagttg ataagttaga tttagattag 600 gaggaattta aacatgtatg agggaaacaa catgcgttct atgatgggta catcatatga 660 agattcaaga ttaaataaac gaacagaatt aaatgaaaac atgtcaattg atacaaataa 720 aagtqaaqat aqttatqqtq tacaaattca ttcactttca aaacaatcat ttacaggtga 780 cgttgaggag gaataataaa ttatggcaca acaatctaca aaaaatgaaa ctgcactttt 840 agtagcaaag tcagctaaat cagcgttaca agattttaat catgattatt caaaatcttg 900 gacatttggc gacaaatggg ataattcaaa tacaatgttc gaaacatttg taaataaata 960 tttattccct aagattaatg agactttatt aatcgatatt gcattaggta atcgttttaa 1020 ttggttagct aaagagcaag attitattgg acaatatagt gaagaatacg tgattatgga 1080 cacagtacca attaacatgg acttatctaa aaatgaggaa ttaatgttga aacgtaatta 1140 tccacgtatg gcaactaagt tatatggtaa cggaattgtg aagaaacaaa aattcacatt 1200 aaacaacaat gatacacgtt tcaatttcca aacattagca gacgcaacta attacgcttt 1260 aggtgtatac aaaaagaaaa tttctgatat taatgtatta gaagaaaaag aaatgcgtgc 1320 aatgttagtt gattactcat tgaatcaatt atccgaaaca aatgtacgta aagcaacatc 1380 aaaagaagat ttagcaagca aagtttttga agcaatccta aacttacaaa acaacagtgc 1440 taaatataat gaagtacatc gtgcatcagg tggtgcaatt ggacaatata caactgtatc 1500 aaaattaaaa gatattgtga ttttaacaac agattcatta aaatcttatc ttttagatac 1560 taagattgca aacacattcc agattgcagg cattgatttc acagatcacg ttattagttt 1620 tgacgactta ggtggcgtgt ttaaagtaac aaaagaattt aagttacaaa accaagattc 1680 aattgacttt ttacgtgcgt atggagatta tcaatcacaa ttaggagata caattccagt 1740 tggtgctgta tttacttatg atgtatctaa acttaaagag tttactggca acgttgaaga 1800 aattaaacca aaatcagatt tatatgcgtt tattttggat attaattcaa ttaaatataa 1860 acgttacaca aaaggtatgt taaaaccacc attccataac cctgaatttg atgaagttac 1920 acactggatt cattactatt catttaaagc cattagtcca ttctttaata aaattttaat 1980 tactgaccaa gatgtaaatc caaaaccaga ggaagaatta caagaataaa aggagcgtaa 2040 aatatgaaca acgataaaag aggtttaaac gttgagttat caaaggaaat cagcaaaaga 2100 gttgttgaac atcgcaacag atttaaacgt cttatgttta atcgttattt ggaattttta 2160 ccgctactaa tcaactatac caatcgtgat acggttggta tagattttat tcagttagaa 2220 tcagctttaa gacaaaacat taatgtagtt gttggtgaag ctagaaataa gcaaattatg 2280 attettggtt atgtaaataa caettaettt aateaageae caaattttte ateaaaettt 2340 aatttccaat ttcaaaaacg attaactaaa gaagatatat attttattgt acctgactat 2400

ttaatacctg atgattgtct acaaattcat aagctatatg ataactgtat gagtggtaac 2460 tttgttgtca tgcaaaataa accaattcaa tataatagtg atatagaaat tatagaacat 2520 tatactgatg aattagcaga agttgcttta tctcgctttt ctttaatcat gcaagcaaaa 2580 tttagcaaga tatttaaatc agaaattaat gacgagtcaa tcaatcaact tgtgtccgaa 2640 atatataacg gtgcaccatt tgttaaaatg tcacctatgt ttaatgcaga tgacgatatc 2700 attgatttaa caagtaatag cgtaatccca gcattaactg aaatgaaacg ggaatatcaa 2760 aacaaaatta gtgaattaag taactattta ggcattaatt cattagccgt tgataaagaa 2820 agcggtgttt cagacgaaga ggcaaaaagt aatcgtggat ttaccacatc aaacagtaat 2880 atctatttaa aaggtcgtga accaattacg tttttatcaa agcgttatgg tttagatatt 2940 aaaccgtatt acgatgatga aacaacgtct aaaatatcaa tggtagacac actttttaaa 3000 gatgaaagca gtgatataaa tggctagata cacaatgact ttatacgatt tcattaaatc 3060 agaattgatt aaaaaaggtt tcaatgaatt tgtaaatgat aataaattaa cgttttatga 3120 tgatgaattt caattcatgc aaaaaatgct gaagttcgac aaagacgttt tagctatcgt 3180 taatgaaaaa gtatttaaag gtttttcatt gaaagatgaa ttatcagatt tactttttaa 3240 aaaatcattt acgattcatt ttttagatag agaaatcaac agacaaacag ttgaagcatt 3300 tggcatgcaa gtgattactg tatgtattac acatgaggat tatttaaatg tggtttattc 3360 atcaagtgaa gttgaaaaat acttacaatc acaaggcttc acagaacaca atgaagatac 3420 aacaagtaac actgatgaaa catcgaatca aaatgctaca tctttagaca attcaactgg 3480 catgactgca aacagaaacg cttatgtgtc attaccacaa agtgaggtta acattgatgt 3540 tgataataca acgttacgat tcgctgataa taatacgatt gataacggta aaactgtgaa 3600 taaatcgagt aacgaaagta atcaaaacgc aaaacgtaat caaaatcaaa aaggtaatgc 3660 aaaaggtaca caattcacta agcagtattt aattgataat attgataaag cgtacgattt 3720 aagaaagaaa attttaaatg aatttgataa aaaatgtttt ttacaaattt ggtagaggtg 3780 gttaaataat ggcatataat gaaaacgatt ttaaatattt tgatgacatt cgtccatttt 3840 tagacgaaat ttataaaacg agagaacgtt atacaccgtt ttacgatgat agagcagatt 3900 ataatactaa ttcaaaatca tattatgatt atatttcaag attatcaaaa ctaattgaag 3960 tattagcacg tcgtatttgg gactatgaca atgaattaaa aaaacgtttc aaaaattggg 4020 acgacttaat gaaagcattt ccagagcaag cgaaagactt atttagaggt tggttaaacg 4080 acggtacgat tgacagtatt attcatgacg agtttaaaaa atatagcgca ggattaacat 4140 cggcatttgc tttatttaaa gttactgaaa tgaaacaaat gaatgacttt aaatcagaag 4200 ttaaagactt aattaaagat attgaccgtt tcgttaatgg gtttgaatta aatgagcttg 4260 aaccaaagtt tgtgatgggc tttggtggta ttcgcaacgc agttaaccaa tctattaata 4320 ttgataaaga aacaaatcac atgtactcta cacaatccga ttctcaaaaa cctgaaggtt 4380 tttggataaa taaattaaca cctagtggtg acttaatttc aagcatgcgt attgtacagg 4440 gtggtcatgg tacaacaatc ggattagaac gtcaatccaa tggtgaaatg aaaatctggt 4500 tacatcacga tggtgttgca aaactgttac aagtcgcata taaagataat tatgtattag 4560 atttagaaga ggctaaaggt ttaacagatt atacaccaca gtcactttta aacaaacaca 4620 catttacacc gttaattgat gaagcaaatg acaaactcat tttaagattc ggtgacggaa 4680 caatacaggt tcgttcaaga gcagacgtaa aaaatcacat tgataatgta gaaaaagaaa 4740 tgacaattga taattcagaa aacaatgata atcgttggat gcaaggcatt gctgttgatg 4800 gtgatgattt atactggtta agtggtaaca gttcagttaa ttcacatgtt caaatcggta 4860 aatattcatt aacaacaggt caaaagattt atgattatcc atttaagtta tcatatcaag 4920 acggtattaa tttcccacgt gataacttta aagagcctga gggtatttgc atttatacaa 4980 atccaaaaac aaaacgtaaa tcgttattac ttgctatgac aaacggcggt ggtggaaaac 5040 gtttccataa tttatatggt ttcttccaac ttggtgagta tgaacacttt gaagcattac 5100 gcgcaagagg ttcacaaaac tataaattaa caaaagacga cggtcgtgca ttatctattc 5160 cagaccatat cgacgattta aatgacttaa cgcaagctgg tttttattat attgacgggg 5220 gtactgcaga aaaacttaag aatatgccaa tgaatggtag caagcgtata attgacgctg 5280 gttgtttcat taatgtatac cctacaacac aaacattagg tacggttcaa gaattaacac 5340 gtttctcaac aggtcgtaaa atggttaaaa tggtgcgtgg tatgacttta gacgtattta 5400 cgttaaaatg ggattatgga ttatggacaa caatcaaaac tgacgcacca tatcaagaat 5460 atttggaagc aagtcaatac aataactgga ttgcttatgt aacaacagct ggtgagtatt 5520

acattacagg taaccaaatg gaattattta gagacgcgcc agaagaaatt aaaaaagtgg 5580 gtgcatggtt acgtgtgtca agtggtaacg cagtcggtga agtaagacaa acattagagg 5640 ctaatatatc ggaatataaa gaattcttca gtaatgttaa tgcggaaaca aaacatcgtg 5700 aatatggttg ggtagcaaaa catcaaaaat aggagtgata taaatgaaat cacaacaaca 5760 agcaaaagaa tggatatata agcatgaggg ggcaggtgtt gactttgatg gtgcatatgg 5820 atttcaatgt atggacttat cagttgctta tgtgtattac attactgacg gtaaagttcg 5880 catgtggggt aatgctaaag acgcgataaa taatgacttt aaaggtttag cgacggtgta 5940 taaaaataca ccgagcttta aacctcaatt aggggacgtt gctgtatata caaatggaca 6000 atatggacat attcaatgtg tgttaagtgg aaatcttgat tattatacat gcttagaaca 6060 aaactggtta ggcggcggtt ttgacggttg ggaaaaagca accattagaa cacattatta 6120 tgacggtgta actcacttta ttagacctaa attttcaggt agtaatagca aagcattaga 6180 aaatgaaaat ggtacattta catgtggttt tttaccaata tttgcacgtg tcggtagtcc 6300 aaaattatca gaacctaatg gctattggtt ccaaccaaac ggttatacac catataacga 6360 agtttgttta tcagatggtt acgtatggat tggttataac tggcaaggca cacgttatta 6420 tttaccagtg cgccaatgga atggaaaaac aggtaatagt tacagtgttg gtattccttg 6480 gggggtgttc tcataatggg tattttagcc tttttctttg aatttagttg gaaaagatac 6540 aaataagagg tgtaaacaat ggctgataga atcgtaagaa gtttaagaca agttgaaaca 6600 attgaacgtt tattggagga aaaaaatgag aaagttaacg aattttaagt ttttctataa 6660 cacaccgttt acagactatc aaaacacgat tcattttaat agtaataaag aacgtgatga 6720 ttatttttta aatggtcgtc attttaaatc gttagactat tcaaaacaac cgtataattt 6780 tatacgtgat agaatggaaa tcaatgttga tatgcagtgg catgacgcac aaggtattaa 6840 ctacatgacg tttttatcag attttgagga tagaagatat tacgcttttg taaaccaaat 6900 cgaatacgtg aatgacgttg tggttaaaat atattttgtc attgatacca ttatgacgta 6960 tacacaaggg aatgtattag agcaactctc aaacgtcaat attgaacgtc aacatttatc 7020 aaaacgcacg tataactata tgttaccaat gttacgtaat aatgatgatg tgttaaaagt 7080

atcaaataaa aactatgttt ataaccaaat gcaacaatat ttggaaaatt tagtattatt 7140 ccagtcaagc gctgatttat caaagaaatt tggtactaaa aaagagccaa acttagatac 7200 gtcaaaaggt acgatttatg acaatatcac atcaccagtc aacttatacg ttatggaata 7260 tggtgacttt attaacttta tggataaaat gagtgcctat ccatggatta cgcaaaactt 7320 tcaaaaggtt caaatgttac ctaaagactt tattaataca aaagacttag aggacgttaa 7380 aaccagtgaa aaaattacag gattaaaaac attaaaacag ggtggtaaat caaaagaatg 7440 gagtctaaaa gatttatcat taagtttctc aaatcttcaa gagatgatgt tatctaaaaa 7500 agatgaattt aaacatatga tacgtaatga gtatatgaca attgaatttt atgactggaa 7560 tggaaatacg atgttactcg acgctggtaa gatttcacaa aaaactggtg ttaagttacg 7620 tacaaaatca attattggtt atcataatga agttcgagta tatccagtag attataacag 7680 tgctgaaaac gacagaccaa tactcgctaa aaataaagaa atattgattg atacgggttc 7740 attottaaat acaaatataa catttaatag ttttgcacaa gtaccaatat taatcaataa 7800 tggtatctta ggacaatcac aacaagccaa ccgacaaaaa aatgcagaaa gtcaattaat 7860 tacaaatcgt attgataatg tattaaatgg tagcgacccg aaatcacgct tttatgacgc 7920 tgtgagtgta gcaagtaatt taagtccaac tgctttattt ggtaagttta atgaagaata 7980 taatttctac aaacaacaac aagctgaata taaagattta gccttacaac caccttctgt 8040 aactgaatca gaaatgggca acgcattcca aattgcgaat agcattaacg gtttaacgat 8100 gaaaattagt gtaccgtcac ctaaagaaat tacattttta caaaaatatt atatgttgtt 8160 tggttttgaa gtgaatgact ataattcatt tattgaacca attaacagta tgactgtttg 8220 caattattta aaatgtacag gtacgtatac tatacgtgac atcgacccca tgttaatgga 8280 acaattaaaa gcaattttag aatctggtgt aagattttgg cataatgacg gttcaggtaa 8340 tccaatgtta caaaatccat taaataacaa atttagagag ggggtataat atgaacgaag 8400 taaaattcag atttacagac tcagaagcgt ttcacatgtt tatatacgct ggggatttaa 8460 aattactcta ctttttattt gtattaatgt tcgttgatat tattacaggt atttcaaaag 8520 caattaaaaa taataactta tggtcaaaaa aatcaatgag aggattttct aaaaaattat 8580 tgatattctg tattatcatt ttagcaaaca tcattgacca gattttacaa ttaaaaggtg 8640 gtctactcat gattacaata ttttattata ttgcaaatga gggactttct attgtagaaa 8700 attgtgcaga aatggacgta ttagtaccag aacaaattaa agataaatta agagtcatta 8760 aaaatgatac tgaaaagagt gataacaatg aacgatcaag agaagataga taaatttacg 8820 cattcctata ttaatgatga ttttggttta acgatagacc agttagtccc taaagtaaaa 8880 ggatatgggc gctttaatgt atggcttggt ggtaatgaaa gtaaaatcag acaagtatta 8940 aaagcagtaa aagagatagg tgtttcacct actctttttg ccgtatatga aaaaaatgag 9000 ggttttagtt ctggacttgg ttggttaaac catacgtctg cacgtggtga ttatttaaca 9060 gatgctaaat tcatagcaag aaagttagta tcacaatcaa aacaagctgg acaaccgtct 9120 tggtatgacg caggtaacat cgtccacttt gtaccacaag acgtacaaag aaaaggtaat 9180 gcagattttg caaaaaatat gaaagcaggt acaattggac gtgcatatat tccattaaca 9240 gcagctgcta cttgggcggc atattatcct ttaggtttga aagcatcata taacaaagta 9300 caaaactatg gtaatccatt tttagacggt gcgaatacta ttctagcttg gggtggtaaa 9360 ttagacggta aaggtggatc acctagtgat tcgtctgaca gtggtagtag tggtgacagt 9420 ggtagttcac tactcgcttt agcaaaacaa gccatgcaag aattattaaa aaaaatacaa 9480 gacgcattac aatgggacgt tcatagtatt ggtagtgata aattttttag taatgattat 9540 tttacattag aaaaaacatt taacaacaca tatcatatta aaatgacgat tggtttactt 9600 gattcattaa aaaaactgat tgatagcgtt caagtagata gtgggagtag tagttctaat 9660 cctactgatg atgacggaga ccataaacca attagtggta aatcagtcaa gccaaatgga 9720 aaaagtggtc gtgtgattgg tggtaactgg acatatgcac agttaccaga aaaatataaa 9780 aaagcaattg gtgtaccttt attcaaaaaa gaatacttat acaaaccagg taacatattt 9840 cctcaaacgg gtaatgcagg acaatgtaca gaattaacat gggcgtatat gtcacaacta 9900 catggtaaaa gacaacctac cgacgacggt caaataacaa acggtcagcg tgtatggtac 9960 gtctataaaa agttaggtgc aaaaacaaca cataatccaa cagtaggtta tggtttctct10020 agtaaaccac catacttaca agcaactgca tatggtattg gtcacacagg tgttgttgta10080 gcagtttttg aagatggttc gtttttagtt gcaaactata atgtaccacc atatgttgca10140 ccatcacgtg tggtattgta tacactcatt aatggcgtac caaataatgc tggtgataat10200

attgtattct ttagtggtat tgcttaatta actatgctat aatgaacaca tgctagtaat10260 gctagtaaat aaaatacaaa acataatcaa ttttcgtaca catttttcat gttatctcaa10320 aaagaaaagg agactgttat tttaacagtt gcctttttt atttcatcat gttcacgttt10380 ${\tt taatatatgc}\ \ {\tt aaatcagatt}\ \ {\tt tgttatgtac}\ \ {\tt tgaacgttca}\ \ {\tt actggaaata}\ \ {\tt agtcgttaag10440}$ tgaaaatgaa ccgatgtcac tttcaatata aagaatatca tcaaattgac tatggtcgaa10500 attttctcta gcgtctttta atataaattc acgtttcata ttaagttcat cagtaaaata10560 ttcatcatat acattaccac atacaatttc agttttagac ggatatatcg atattgtacc10620 ttgctcatta tagatacttt tattgttttc aataatggca ccgtcaaaga attgttcacg10680 tacaaaggtt tcaaaatcga cgcttgtatc aaaggcgttt ttcggtatac cagcagaagc10740 aattttaatc tttccattca cttcatatgc atatttctta tgattcagta caaacatctt10800 atctatctgt tcgttttcaa tatcccattt acctaaggct atcgggtcga ataaactggg10860 gttcaataag ggtttaacaa cggatttcat atacaaacta tcagtatcgc aataaataaa10920 attgtcgtca atttcacttt ccgttaagta ttggaaagga accaataagt tatacaatga10980 acgtgatgtg acaaatgtag agaataatat attacgttca gtgtttttgt aaccgttaat11040 gatattgtat agttcattgt tatcatctaa acggaataag ttaaaatgtg aacgtaatgc11100 aggtatgcca tataatccat ttaaaacgac tttagataac ataacctcct catttgagta11160 tgggtgttcg ttgatatcat cagtaatgtg atagtcgtaa ggtgatgtca tattgatttt11220 gttttttaac ttaccttgtg ttttaataaa atagttttga aaaataatat cacgtgcatg11280 aaagtattca cattcatata taacaaacga attaacacgt atatgcatgc aatcaatacc11340 cgtaatgtct tgaatcattc ttaatgtatt tgtattgata ttaacgtaat cattatcatt11400 attatagtat tttacaatca tttgacgtaa tacacgtgat ttaattttaa ttaataaatc11460 atcgttaaat acatctttat caatcttata taatgaaaaa taattgtcat catctaaaaa11520 agtagggatt aacgttggtt ctgaatagtg ttcgtaaaag tataaccatg ttggaatttt11580 ttcatgatac atcacataag gataactcga attgatgtca atagaaaaac aaggctcatc11640 aattagtttg tttatgtatt tggtgttata catatttaaa ccaccacgat agaatgattt11700 aatatagtca taaaaattca tatcatggaa atgataatgt gtataagata ttttaatatc11760

ttgatattgg ttgagtaact gaaaacgtgt catttcatta ttcaagtaag attccataat11820 attcaatgaa aatgttaatt tgttatagtc aaaatttgga aatatatcac tataatgaat11880 atggcacata cctaatataa tcacgtcatt atgaatgtat gtaagttgtt caggtgtgag11940 ttttgcaaaa catttcacag catagtcata ggcttcacta tcattcatat cattatcttt12000 atcaaaaatc gtataattaa aatctgtttt aagttgtgat tctgttaaat aaccaccatc12060 aagtaatttc ttacctaatg ttgcaattga tgtattggtt ttcataaagt tatcaataat12120 attaaattta aaaccattta aaaacattgt taaatctaaa ttgattgaag atttaacacg12180 tttttctaaa attacatttt gatttttggc taaaatagta gcctctttca tttttaatgt12240 gtgttcattt tcttctgcag attttaaata tatattttcg cgtgtaatat tatcaaaata12300 acgcatggtg tctttaagta aaaaatgatt atcgtattta ttacagttat gtgcaatcat12360 gataatatct gtttttgatt ttgtgattgt atcacgtctt ttcacatacg tataaaatgc12420 gtcataaaaa gattcgaaac tcggaaatac ttcaacatca atttcataac cattaaacca12480 accaattgct acagaataag taacgttttt atatttggtt ggtttttttc gtccgttaac12540 tttattgtac gctaatgttt ctatatccca gtataaaatc attcgacgtt catgtttatg12600 atattgcatg cattctagta atcccataat cttacacacc ttttataagc catattgttt12660 cattagatac tttttcgtat tctctatata gttatcttcg tatatttttt cttttctttc12720 aaactcactc atatttttct tcatttcatt ttttatatga aattttataa ttttattcat12780 atctaaatat aaatatctat cattatcaac cacgtaattt ttagagtaag cattgtcaaa12840 atgtaaattg cttggattgt agtaataacg ttccatgttt tctttataaa acatatcatc12900 acgtaaatag gtaacatgat tgtctatatc cctaatttta gtacaaaatt catattgttt12960 tgtatatggt acaacgataa tatttgtcat aaaagtagtt acattataca tgactttaat13020 atatttatca tcagttttga tatagaagaa atcaccgttt tgattgatgt gatttcttaa13080 attatcatcc gccaaattat attcgttaaa ttcaaattct ccagttgtca tagcgtcgtc13140 atttgaatta aacgcacgtg tgttacgttt ttcattcacg taatcgtttc gtcgcatttc13200 taaaaaaatg tttttgtaaa gtcttgatgt attcatttta tgcttttgta ataaattgta13260 tatatttaaa ttggataata taggacttga aaagttgact gcattaccta gtaaaaacat13320 tttagggaat ccaatataat caacgttacc atggttacgg tcgattgatt catatattgt13380 ttttaactta tcccactcat caattaaata atcatcttca agtgctaaaa actcatcata13440 tataataata ggatagtgtt ttaaaaagtt agaatgatat tttaaatcag tggcactatt13500 caaatctgta atcacaccaa tttctttatc ttgatagata atagctaaat agtccctagc13560 acttetgaac gtgacacgtt ttgatttaaa tagtggattt teatetatga tttetteaat13620 aaaatcacgg taagcgtcac gtaatgtata atgacgtgat aataaagtaa attttatatc13680 aagtttaata gctaaataaa taaaaaatga aacatagttg aacgattttc catcagaacg13740 gtttgaaata gatatataat aatctatatc atcattcata agttcatcaa ctaattctat13800 ttgattatac ttatctggga ttttttttct gacatgattg acagcatttt gataatctct13860 taccatgtct aaacgatttt gttttaccat gtttttgctc cttgtaatag tttatgatgt13920 cgtttacagt gttaaattta ttcgtcaaat gttgcataat ataaaaagtt atacctcaca13980 tetteateat caatatttgt caetggteta tetgatttae caatttettt atataaagta14040 tegatttett taatatattt atacattgaa gaattattat ttttagettg taaattatat14100 aaagcgtatt tatgcttttt agcgttttta ttattagaat catcattacg gttatatatt14160 tcaagaatat aatttaattt tttatgtctt gaacctctta ccaatgatac agcatttaca14220 tatgatacgt ttctttcttt aggaaaatag ggcagatgtg caaaatgttt ccatgtgtca14280 atgtacgcct cttgtaaatc tttatcatca aatttaaaat taacattact aaaatcattt14340 aaaaataaat ctttttcttg ctcttttcta gcttctcttt ctttttcca tctatccatt14400 tcagacgtat gtctaaccaa tgttatcaac ctccatataa agcataaata accattaaaa14460 agataatata gaatataatc aatgtagtga ataaaacacc aaatgacacg cgtatatgca14520 gtgtcataag tatgataagt gtaattaaaa atgctaaaag gaaaacaatg gctatgttta14580 ataggttatt catggtcaat cactttccca ttatcgtata tgactttgtt ttgataaata14640 atcattaatt cgctttcaag aggtttatca aaatttgata atacgtcgtc aattgtaacg14700 tttaataaaa tttctcttat taattcatta cttaaataat ttctataata aaatacaagt14760 atattaaaaa catgtttttt aatatcaatg tcgatatcta acgtaaataa ctctttttca14820 atttcaaaat catcatattg tttgtcaaac tcaatataca catcacccat atttattttt14880

actatacatt ttttattaga tgaagtaaat ttttcaaatt tatcattata ataatctcta14940 tttgttaaaa ggtaataaat taaattattt aatctaaaag tagttttaat tttcattttt15000 atatctcctt aatgtattct atgatatacg cgtatttttt agtgaacagg ttatattcat15060 aatatgaata tacaacttta gcgtcatata aatcttcaaa cattgagatt tgatgtggaa15120 aatgtccttt aatctcatcg caatataata ataccgtttt gtatttacgt tccatttaaa15180 cacctcataa aaaatagggg ataagtatcc cctatgaaat tgtattaaaa tgatacttga15240 ccaaaattga ttgagtaacc tttttgacct tttttgtttt catattcata aattgtgaat15300 tgaacttctc cagcattgat aatgtcaaca acgtcctcat ctgctctcat ttctttaatt15360 aattctgtta agtggttcgg taagtttacg ttatagtcat cagtgacgat aacaccttgt15420 tcaccgaatt ttgattcttt gtttgtgaat aatgctctaa cgatatactc ttttttcata15480 ccgtattttt ctactaattc tgatagtttg ataaattctc tttctttttc ctcaaattca15540 aatctcgcta atgtgttttg gtgtcttgat aaaatatctt ttacgtttgt cattttattt15600 ctcctcttat ttaaattatt tgctttctgc aattgcgatt tgtagtaaat cattgtaata15660 aacttgaatt gttttcgttg tgcgtgtagt ggacaatagt ttacatgtgt ctggtaataa15720 ttcttttgct tgtgttttgg ttaaatgata ctcgtgaagt ggtaaaaatt cctcaatgta15780 ttcattatca tcatctaagt aatgaagtat ataacctttg acacgtaagg taacaatgtc15840 gtcaactttc attattatat cactcctttc taaaaaacgt aaacgttata cgtttcataa15900 aatcctttat gcatattcca ttgttctatt gggtcatcac cagcaatata agacaatatt15960 gattctggtt tagtttcgtt gtttagttca tcatttaaga attgaacaac agaactatta16020 tagtttaata atagttgttg gcaagccgat aataagttaa ttgcattgtc aaatgtataa16080 gctggattcc attgaatcag tttattgaat agttgcaaca tttcagtata ggcttgtcct16140 ttttcttctg gtgcattatc aacattaacc attattatca cttcctaata aagttgaaat16200 tacgcgtaaa acagaattat gatttaaatc ttcaatttca tcaatgtcaa catcataaaa16260 tgaaatttca ttttctgttc tatcaaataa cgctatacat aaacttccat tcttaaaacg16320 aaaaacatgc ttcaactcaa tgttttttgt ttcattttcc atttttgtta ctccttgttt16380 tgattacata cttagtatag caaacgttta aaagttttgt caatagtttt tcttaaaaaa16440

017.176341.1

gtttaaataa ttttaaaact actatttaat agaagaaata agattttaag ttcaaatcat16500 aattttgaat aaaagtcaat agatacataa attttgtatt tgatgaatat gtaataggtt16560 agataagttg gttaagttgt tgcacagtat ttttaagttt agtaaagaaa tgataagtaa16620 atttataagt tttgatttgt ataatcgttt attttaaacc ggtggggt 16668

<210> 2

<211> 2286

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 2

60 atgggattac tagaatgcat gcaatatcat aaacatgaac gtcgaatgat tttatactgg gatatagaaa cattagcgta caataaagtt aacggacgaa aaaaaccaac caaatataaa 120 180 aacgttactt attctgtagc aattggttgg tttaatggtt atgaaattga tgttgaagta 240 tttccgagtt tcgaatcttt ttatgacgca ttttatacgt atgtgaaaag acgtgataca atcacaaaat caaaaacaga tattatcatg attgcacata actgtaataa atacgataat 300 360 cattttttac ttaaagacac catgcgttat tttgataata ttacacgcga aaatatatat ttaaaatctg cagaagaaaa tgaacacaca ttaaaaatga aagaggctac tattttagcc 420 480 aaaaatcaaa atgtaatttt agaaaaacgt gttaaatctt caatcaattt agatttaaca atgtttttaa atggttttaa atttaatatt attgataact ttatgaaaac caatacatca 540 attgcaacat taggtaagaa attacttgat ggtggttatt taacagaatc acaacttaaa 600 660 acagatttta attatacgat ttttgataaa gataatgata tgaatgatag tgaagcctat 720 gactatgctg tgaaatgttt tgcaaaactc acacctgaac aacttacata cattcataat gacgtgatta tattaggtat gtgccatatt cattatagtg atatatttcc aaattttgac 780 840 tataacaaat taacattttc attgaatatt atggaatctt acttgaataa tgaaatgaca cgttttcagt tactcaacca atatcaagat attaaaatat cttatacaca ttatcatttc 900 960 catgatatga attittatga ctatattaaa tcattctatc gtggtggttt aaatatgtat aacaccaaat acataaacaa actaattgat gagccttgtt tttctattga catcaattcg 1020 agttatcctt atgtgatgta tcatgaaaaa attccaacat ggttatactt ttacgaacac 1080

tattcagaac caacgttaat ccctactttt ttagatgatg acaattattt ttcattatat 1140 aagattgata aagatgtatt taacgatgat ttattaatta aaattaaatc acgtgtatta 1200 cgtcaaatga ttgtaaaata ctataataat gataatgatt acgttaatat caatacaaat 1260 acattaaqaa tqattcaaqa cattacqqqt attqattqca tqcatatacq tqttaattcq 1320 tttgttatat atgaatgtga atactttcat gcacgtgata ttatttttca aaactatttt 1380 attaaaacac aaggtaagtt aaaaaacaaa atcaatatga catcacctta cgactatcac 1440 attactgatg atatcaacga acacccatac tcaaatgagg aggttatgtt atctaaagtc 1500 qttttaaatg qattatatgg catacetgca ttacgttcae attttaactt attecgttta 1560 gatgataaca atgaactata caatatcatt aacggttaca aaaacactga acgtaatata 1620 ttattctcta catttgtcac atcacgttca ttgtataact tattggttcc tttccaatac 1680 ttaacggaaa gtgaaattga cgacaatttt atttattgcg atactgatag tttgtatatg 1740 aaatccgttg ttaaaccctt attgaacccc agtttattcg acccgatagc cttaggtaaa 1800 tgggatattg aaaacgaaca gatagataag atgtttgtac tgaatcataa gaaatatgca 1860 tatqaagtga atggaaagat taaaattgct tctgctggta taccgaaaaa cgcctttgat 1920 acaagcgtcg attttgaaac ctttgtacgt gaacaattct ttgacggtgc cattattgaa 1980 aacaataaaa qtatctataa tgagcaaggt acaatatcga tatatccgtc taaaactgaa 2040 attgtatgtg gtaatgtata tgatgaatat tttactgatg aacttaatat gaaacgtgaa 2100 tttatattaa aagacgctag agaaaatttc gaccatagtc aatttgatga tattctttat 2160 attgaaagtg acatcggttc attttcactt aacgacttat ttccagttga acgttcagta 2220 cataacaaat ctgatttgca tatattaaaa cgtgaacatg atgaaataaa aaaaggcaac 2280 2286 tgttaa

<210> 3

<211> 1944

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 3

atggcatata atgaaaacga ttttaaatat tttgatgaca ttcgtccatt tttagacgaa 60 atttataaaa cgagagaacg ttatacaccg ttttacgatg atagagcaga ttataatact 120

aattcaaaat catattatga ttatatttca agattatcaa aactaattga agtattagca 180 240 cgtcgtattt gggactatga caatgaatta aaaaaacgtt tcaaaaattg ggacgactta 300 atgaaagcat ttccagagca agcgaaagac ttatttagag gttggttaaa cgacggtacg attgacagta ttattcatga cgagtttaaa aaatatagcg caggattaac atcggcattt 360 gctttattta aagttactga aatgaaacaa atgaatgact ttaaatcaga agttaaagac 420 480 ttaattaaag atattgaccg tttcgttaat gggtttgaat taaatgagct tgaaccaaag 540 tttgtgatgg gctttggtgg tattcgcaac gcagttaacc aatctattaa tattgataaa gaaacaaatc acatgtactc tacacaatcc gattctcaaa aacctgaagg tttttggata 600 aataaattaa cacctagtgg tgacttaatt tcaagcatgc gtattgtaca gggtggtcat 660 720 ggtacaacaa tcggattaga acgtcaatcc aatggtgaaa tgaaaatctg gttacatcac gatggtgttg caaaactgtt acaagtcgca tataaagata attatgtatt agatttagaa 780 gaggctaaag gtttaacaga ttatacacca cagtcacttt taaacaaaca cacatttaca 840 900 ccgttaattg atgaagcaaa tgacaaactc attttaagat tcggtgacgg aacaatacag gttcgttcaa gagcagacgt aaaaaatcac attgataatg tagaaaaaga aatgacaatt 960 gataattcag aaaacaatga taatcgttgg atgcaaggca ttgctgttga tggtgatgat 1020 ttatactggt taagtggtaa cagttcagtt aattcacatg ttcaaatcgg taaatattca 1080 aatttcccac gtgataactt taaagagcct gagggtattt gcatttatac aaatccaaaa 1200 acaaaacgta aatcgttatt acttgctatg acaaacggcg gtggtggaaa acgtttccat 1260 aatttatatg gtttcttcca acttggtgag tatgaacact ttgaagcatt acgcgcaaga 1320 ggttcacaaa actataaatt aacaaaagac gacggtcgtg cattatctat tccagaccat 1380 atcgacgatt taaatgactt aacgcaagct ggtttttatt atattgacgg gggtactgca 1440 gaaaaactta agaatatgcc aatgaatggt agcaagcgta taattgacgc tggttgtttc 1500 attaatgtat accetacaac acaaacatta ggtacggtte aagaattaac acgtttetea 1560 acaggtcgta aaatggttaa aatggtgcgt ggtatgactt tagacgtatt tacgttaaaa 1620 tgggattatg gattatggac aacaatcaaa actgacgcac catatcaaga atatttggaa 1680

017.176341.1

<210> 4

<211> 1764

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 4

atgagaaagt taacgaattt taagtttttc tataacacac cgtttacaga ctatcaaaac 60 120 acgattcatt ttaatagtaa taaagaacgt gatgattatt ttttaaatgg tcgtcatttt aaatcgttag actattcaaa acaaccgtat aattttatac gtgatagaat ggaaatcaat 180 gttgatatgc agtggcatga cgcacaaggt attaactaca tgacgttttt atcagatttt 240 300 gaggatagaa gatattacgc ttttgtaaac caaatcgaat acgtgaatga cgttgtggtt aaaatatatt ttgtcattga taccattatg acgtatacac aagggaatgt attagagcaa 360 420 ctctcaaacg tcaatattga acgtcaacat ttatcaaaac gcacgtataa ctatatgtta 480 ccaatgttac gtaataatga tgatgtgtta aaagtatcaa ataaaaacta tgtttataac caaatgcaac aatatttgga aaatttagta ttattccagt caagcgctga tttatcaaag 540 600 aaatttggta ctaaaaaaga gccaaactta gatacgtcaa aaggtacgat ttatgacaat 660 atcacatcac cagtcaactt atacgttatg gaatatggtg actttattaa ctttatggat 720 aaaatgagtg cctatccatg gattacgcaa aactttcaaa aggttcaaat gttacctaaa 780 gactttatta atacaaaaga cttagaggac gttaaaacca gtgaaaaaat tacaggatta aaaacattaa aacagggtgg taaatcaaaa gaatggagtc taaaagattt atcattaagt 840 900 ttctcaaatc ttcaagagat gatgttatct aaaaaagatg aatttaaaca tatgatacgt 960 ggtaagattt cacaaaaaac tggtgttaag ttacgtacaa aatcaattat tggttatcat 1020 aatgaagttc gagtatatcc agtagattat aacagtgctg aaaacgacag accaatactc 1080 gctaaaaata aagaaatatt gattgatacg ggttcattct taaatacaaa tataacattt 1140 aatagttttg cacaagtacc aatattaatc aataatggta tcttaggaca atcacaacaa 1200 gccaaccgac aaaaaaatgc agaaagtcaa ttaattacaa atcgtattga taatgtatta 1260 aatggtagcg acccgaaatc acgcttttat gacgctgtga gtgtagcaag taatttaagt 1320 ccaactgctt tatttggtaa gtttaatgaa gaatataatt tctacaaaca acaacaagct 1380 gaatataaag atttagcctt acaaccacct tctgtaactg aatcagaaat gggcaacgca 1440 ttccaaattg cgaatagcat taacggttta acgatgaaaa ttagtgtacc gtcacctaaa 1500 gaaattacat ttttacaaaa atattatatg ttgtttggtt ttgaagtgaa tgactataat 1560 tcatttattg aaccaattaa cagtatgact gtttgcaatt atttaaaatg tcacggtca 1620 tatactatac gtgacatcga ccccatgtta atggaacaat taaaagcaat tttagaatct 1680 ggtgtaagat tttggcataa tgacggtca ggtaatccaa tgttacaaaa tccattaaat 1740 aacaaattta gagaggggt ataa

<210> 5

<211> 1464

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 5

atgatactga aaagagtgat aacaatgaac gatcaagaga agatagataa atttacgcat 60 tcctatatta atgatgattt tggtttaacg atagaccagt tagtccctaa agtaaaagga 120 tatgggcgct ttaatgtatg gcttggtggt aatgaaagta aaatcagaca agtattaaaa 180 gcagtaaaag agataggtgt ttcacctact ctttttgccg tatatgaaaa aaatgagggt 240 tttagttctg gacttggttg gttaaaccat acgtctgcac gtggtgatta tttaacagat 300 gctaaattca tagcaagaaa gttagtatca caatcaaaac aagctggaca accgtcttgg 360 420 tatgacgcag gtaacatcgt ccactttgta ccacaagacg tacaaagaaa aggtaatgca gattttgcaa aaaatatgaa agcaggtaca attggacgtg catatattcc attaacagca 480 540 gctgctactt gggcggcata ttatccttta ggtttgaaag catcatataa caaagtacaa 600 aactatggta atccattttt agacggtgcg aatactattc tagcttgggg tggtaaatta

gacggtaaag gtggatcacc tagtgattcg tctgacagtg gtagtagtgg tgacagtggt 660 720 agttcactac tcgctttagc aaaacaagcc atgcaagaat tattaaaaaa aatacaagac gcattacaat gggacgttca tagtattggt agtgataaat tttttagtaa tgattatttt 780 acattagaaa aaacatttaa caacacatat catattaaaa tgacgattgg tttacttgat 840 tcattaaaaa aactgattga tagcgttcaa gtagatagtg ggagtagtag ttctaatcct 900 actgatgatg acggagacca taaaccaatt agtggtaaat cagtcaagcc aaatggaaaa 960 agtggtcgtg tgattggtgg taactggaca tatgcacagt taccagaaaa atataaaaaa 1020 gcaattggtg tacctttatt caaaaaagaa tacttataca aaccaggtaa catatttcct 1080 caaacgggta atgcaggaca atgtacagaa ttaacatggg cgtatatgtc acaactacat 1140 ggtaaaagac aacctaccga cgacggtcaa ataacaaacg gtcagcgtgt atggtacgtc 1200 tataaaaagt taggtgcaaa aacaacacat aatccaacag taggttatgg tttctctagt 1260 aaaccaccat acttacaagc aactgcatat ggtattggtc acacaggtgt tgttgtagca 1320 gtttttgaag atggttcgtt tttagttgca aactataatg taccaccata tgttgcacca 1380 tcacgtgtgg tattgtatac actcattaat ggcgtaccaa ataatgctgg tgataatatt 1440 1464 gtattcttta gtggtattgc ttaa

<210> 6

<211> 1248

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 6

atggtaaaac aaaatcgttt agacatggta agagattatc aaaatgctgt caatcatgtc 60
agaaaaaaaa tcccagataa gtataatcaa atagaattag ttgatgaact tatgaatgat 120
gatatagatt attatatac tatttcaaac cgttctgatg gaaaatcgtt caactatgtt 180
tcattttta tttatttagc tattaaactt gatataaaat ttactttatt atcacgtcat 240
tatacattac gtgacgctta ccgtgatttt attgaagaaa tcatagatga aaatccacta 300
tttaaatcaa aacgtgtcac gttcagaagt gctagggact atttagctat tatctatcaa 360
gataaagaaa ttggtgtgat tacagatttg aatagtgcca ctgatttaaa atatcattct 420

480 aactttttaa aacactatcc tattattata tatgatgagt ttttagcact tgaagatgat 540 tatttaattq atgagtggga taagttaaaa acaatatatg aatcaatcga ccgtaaccat 600 ggtaacgttg attatattgg attccctaaa atgtttttac taggtaatgc agtcaacttt 660 tcaagtccta tattatccaa tttaaatata tacaatttat tacaaaagca taaaatgaat acatcaagac tttacaaaaa catttttta gaaatgcgac gaaacgatta cgtgaatgaa 720 780 aaacgtaaca cacgtgcgtt taattcaaat gacgacgcta tgacaactgg agaatttgaa 840 tttaacgaat ataatttggc ggatgataat ttaagaaatc acatcaatca aaacggtgat 900 ttcttctata tcaaaactga tgataaatat attaaagtca tgtataatgt aactactttt atgacaaata ttatcgttgt accatataca aaacaatatg aattttgtac taaaattagg 960 gatatagaca atcatgttac ctatttacgt gatgatatgt tttataaaga aaacatggaa 1020 cgttattact acaatccaag caatttacat tttgacaatg cttactctaa aaattacgtg 1080 gttgataatg atagatattt atatttagat atgaataaaa ttataaaaatt tcatataaaa 1140 1248 tatatagaga atacgaaaaa gtatctaatg aaacaatatg gcttataa

<210> 7

<211> 1227

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 7

60 atggcacaac aatctacaaa aaatgaaact gcacttttag tagcaaagtc agctaaatca 120 gcgttacaag attttaatca tgattattca aaatcttgga catttggcga caaatgggat aattcaaata caatgttcga aacatttgta aataaatatt tattccctaa gattaatgag 180 240 actttattaa tcgatattgc attaggtaat cgttttaatt ggttagctaa agagcaagat 300 tttattggac aatatagtga agaatacgtg attatggaca cagtaccaat taacatggac 360 ttatctaaaa atgaggaatt aatgttgaaa cgtaattatc cacgtatggc aactaagtta 420 tatggtaacg gaattgtgaa gaaacaaaaa ttcacattaa acaacaatga tacacgtttc 480 aatttccaaa cattagcaga cgcaactaat tacgctttag gtgtatacaa aaagaaaatt 540 tctgatatta atgtattaga agaaaaagaa atgcgtgcaa tgttagttga ttactcattg

600 aatcaattat ccgaaacaaa tgtacgtaaa gcaacatcaa aagaagattt agcaagcaaa 660 gtttttgaag caatcctaaa cttacaaaac aacagtgcta aatataatga agtacatcgt gcatcaggtg gtgcaattgg acaatataca actgtatcaa aattaaaaga tattgtgatt 720 780 ttaacaacag attcattaaa atcttatctt ttagatacta agattgcaaa cacattccag attgcaggca ttgatttcac agatcacgtt attagttttg acgacttagg tggcgtgttt 840 900 aaagtaacaa aagaatttaa gttacaaaac caagattcaa ttgacttttt acgtgcgtat ggagattatc aatcacaatt aggagataca attccagttg gtgctgtatt tacttatgat 960 gtatctaaac ttaaagagtt tactggcaac gttgaagaaa ttaaaccaaa atcagattta 1020 tatgcgttta ttttggatat taattcaatt aaatataaac gttacacaaa aggtatgtta 1080 aaaccaccat tccataaccc tgaatttgat gaagttacac actggattca ttactattca 1140 tttaaagcca ttagtccatt ctttaataaa attttaatta ctgaccaaga tgtaaatcca 1200 1227 aaaccagagg aagaattaca agaataa

<210> 8

<211> 984

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 8

atgaacaacg ataaaagagg tttaaacgtt gagttatcaa aggaaatcag caaaagagtt 60 gttgaacatc gcaacagatt taaacgtctt atgtttaatc gttatttgga atttttaccg 120 180 ctactaatca actataccaa tcgtgatacg gttggtatag attttattca gttagaatca 240 gctttaagac aaaacattaa tgtagttgtt ggtgaagcta gaaataagca aattatgatt 300 cttggttatg taaataacac ttactttaat caagcaccaa atttttcatc aaactttaat 360 ttccaatttc aaaaacgatt aactaaagaa gatatatatt ttattgtacc tgactattta atacctgatg attgtctaca aattcataag ctatatgata actgtatgag tggtaacttt 420 480 gttgtcatgc aaaataaacc aattcaatat aatagtgata tagaaattat agaacattat 540 actgatgaat tagcagaagt tgctttatct cgcttttctt taatcatgca agcaaaattt 600 agcaagatat ttaaatcaga aattaatgac gagtcaatca atcaacttgt gtccgaaata

tataacggtg	caccatttgt	taaaatgtca	cctatgttta	atgcagatga	cgatatcatt	660
gatttaacaa	gtaatagcgt	aatcccagca	ttaactgaaa	tgaaacggga	atatcaaaac	720
aaaattagtg	aattaagtaa	ctatttaggc	attaattcat	tagccgttga	taaagaaagc	780
ggtgtttcag	acgaagaggc	aaaaagtaat	cgtggattta	ccacatcaaa	cagtaatatc	840
tatttaaaag	gtcgtgaacc	aattacgttt	ttatcaaagc	gttatggttt	agatattaaa	900
ccgtattacg	atgatgaaac	aacgtctaaa	atatcaatgg	tagacacact	ttttaaagat	960
gaaagcagtg	atataaatgg	ctag				984

<210> 9

<211> 756

<212> DNA

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> atggctagat acacaatgac tttatacgat ttcattaaat cagaattgat taaaaaaggt 60 120 ttcaatgaat ttgtaaatga taataaatta acgttttatg atgatgaatt tcaattcatg 180 caaaaaatgc tgaagttcga caaagacgtt ttagctatcg ttaatgaaaa agtatttaaa ggtttttcat tgaaagatga attatcagat ttacttttta aaaaatcatt tacgattcat 240 300 tttttagata gagaaatcaa cagacaaaca gttgaagcat ttggcatgca agtgattact 360 gtatgtatta cacatgagga ttatttaaat gtggtttatt catcaagtga agttgaaaaa 420 tacttacaat cacaaggett cacagaacac aatgaagata caacaagtaa cactgatgaa 480 acatcgaatc aaaatgctac atctttagac aattcaactg gcatgactgc aaacagaaac qcttatqtqt cattaccaca aagtgaggtt aacattgatg ttgataatac aacgttacga 540 600 ttcgctgata ataatacgat tgataacggt aaaactgtga ataaatcgag taacgaaagt 660 aatcaaaacg caaaacgtaa tcaaaatcaa aaaggtaatg caaaaggtac acaattcact 720 aagcagtatt taattgataa tattgataaa gcgtacgatt taagaaagaa aattttaaat 756 gaatttgata aaaaatgttt tttacaaatt tggtag

<210> <211> <212>	10 753 DNA						
<213>	Stap	hylococcus	aureus Bact	teriophage 4	44 AHJD		
<400> atgaaat	10 tcac	aacaacaagc	aaaagaatgg	atatataagc	atgagggggc	aggtgttgac	60
tttgato	ggtg	catatggatt	tcaatgtatg	gacttatcag	ttgcttatgt	gtattacatt	120
actgac	ggta	aagttcgcat	gtggggtaat	gctaaagacg	cgataaataa	tgactttaaa	180
ggtttag	gcga	cggtgtataa	aaatacaccg	agctttaaac	ctcaattagg	ggacgttgct	240
gtatata	acaa	atggacaata	tggacatatt	caatgtgtgt	taagtggaaa	tcttgattat	300
tatacat	tgct	tagaacaaaa	ctggttaggc	ggcggttttg	acggttggga	aaaagcaacc	360
attagaa	acac	attattatga	cggtgtaact	cactttatta	gacctaaatt	ttcaggtagt	420
aatagca	aaag	cattagaaac	atcaaaagtá	aatacatttg	gaaaatggaa	acgaaaccaa	480
tacggca	acat	attatagaaa	tgaaaatggt	acatttacat	gtggttttt	accaatattt	540
gcacgt	gtcg	gtagtccaaa	attatcagaa	cctaatggct	attggttcca	accaaacggt	600
tatacad	ccat	ataacgaagt	ttgtttatca	gatggttacg	tatggattgg	ttataactgg	660
caaggca	acac	gttattattt	accagtgcgc	caatggaatg	gaaaaacagg	taatagttac	720
agtgtt	ggta	ttccttgggg	ggtgttctca	taa			753
<210> <211> <212>	11 483 DNA						
<213>		hylococcus	aureus Bact	teriophage 4	44 AHJD		
<400>	11	atacatataa	2214421242	taassasaa	aaagagaagc	tagaaaagg	60
	_	-				-	
_			-		attttaaatt		120
•	-			_	cacatctgcc		180
aaagaaa	agaa	acgtatcata	tgtaaatgct	gtatcattgg	taagaggttc	aagacataaa	240
aaattaa	aatt	atattcttga	aatatataac	cgtaatgatg	attctaataa	taaaaacgct	300
aaaaag	cata	aatacgcttt	atataattta	caagctaaaa	ataataattc	ttcaatgtat	360
aaatata	atta	aagaaatcga	tactttatat	aaagaaattg	gtaaatcaga	tagaccagtg	420

acaaatattg	atgatgaaga	tgtgaggtat	aactttttat	attatgcaac	atttgacgaa	480
taa						483
<210> 12 <211> 369 <212> DNA <213> Stap	phylococcus	aureus Bact	ceriophage 4	14 AHJD		
<400> 12 atgacaaacg	taaaagatat	tttatcaaga	caccaaaaca	cattagcgag	atttgaattt	60
gaggaaaaag	aaagagaatt	tatcaaacta	tcagaattag	tagaaaaata	cggtatgaaa	120
aaagagtata	tcgttagagc	attattcaca	aacaaagaat	caaaattcgg	tgaacaaggt	180
gttatcgtca	ctgatgacta	taacgtaaac	ttaccgaacc	acttaacaga	attaattaaa	240
gaaatgagag	cagatgagga	cgttgttgac	attatcaatg	ctggagaagt	tcaattcaca	300
atttatgaat	atgaaaacaa	aaaaggtcaa	aaaggttact	caatcaattt	tggtcaagta	360
tcattttaa						369
<210> 13 <211> 423 <212> DNA <213> Stap	phylococcus	aureus Bact	ceriophage 4	44 AHJD		
<400> 13 atgaacgaag	taaaattcag	atttacagac	tcagaagcgt	ttcacatgtt	tatatacgct	60
ggggatttaa	aattactcta	ctttttattt	gtattaatgt	tcgttgatat	tattacaggt	120
atttcaaaag	caattaaaaa	taataactta	tggtcaaaaa	aatcaatgag	aggattttct	180
aaaaaattat	tgatattctg	tattatcatt	ttagcaaaca	tcattgacca	gattttacaa	240
ttaaaaggtg	gtctactcat	gattacaata	ttttattata	ttgcaaatga	gggactttct	300
attgtagaaa	attgtgcaga	aatggacgta	ttagtaccag	aacaaattaa	agataaatta	360
agagtcatta	aaaatgatac	tgaaaagagt	gataacaatg	aacgatcaag	agaagataga	420
taa						423

<210> <211> <212>	14 411 DNA						
<213>	Stap	ohylococcus	aureus Bact	eriophage 4	14 AHJD		
<400> atgaaaa	14 atta	aaactacttt	tagattaaat	aatttaattt	attacctttt	aacaaataga	60
gattatt	ata	atgataaatt	tgaaaaattt	acttcatcta	ataaaaaatg	tatagtaaaa	120
ataaata	tgg	gtgatgtgta	tattgagttt	gacaaacaat	atgatgattt	tgaaattgaa	180
aaagagt	tat	ttacgttaga	tatcgacatt	gatattaaaa	aacatgtttt	taatatactt	240
gtattt	att	atagaaatta	tttaagtaat	gaattaataa	gagaaatttt	attaaacgtt	300
acaatto	gacg	acgtattatc	aaattttgat	aaacctcttg	aaagcgaatt	aatgattatt	360
tatcaaa	aca	aagtcatata	cgataatggg	aaagtgattg	accatgaata	a	411
<210> <211> <212> <213>	15 279 DNA Stap	phylococcus	aureus Bact	ceriophage 4	44 AHJD		
<400> atgaaaa	15 atgg	tacatttaca	tgtggtttt	taccaatatt	tgcacgtgtc	ggtagtccaa	60
aattato	caga	acctaatggc	tattggttcc	aaccaaacgg	ttatacacca	tataacgaag	120
tttgttt	atc	agatggttac	gtatggattg	gttataactg	gcaaggcaca	cgttattatt	180
taccagt	gcg	ccaatggaat	ggaaaaacag	gtaatagtta	cagtgttggt	attccttggg	240
gggtgtt	ctc	ataatgggta	ttttagcctt	tttctttga .			279
<210> <211> <212> <213>	16 243 DNA Stap	phylococcus	aureus Bact	ceriophage 4	14 AHJD		
<400> gtgacga	16 ataa	caccttgttc	accqaatttt	gattctttgt	ttgtgaataa	tgctctaacg	60
		ttttcatacc					120
			_	_			
cctttt	cct	caaattcaaa	ictcgctaat	gtgttttggt	gtcttgataa	aalatCtttt	180

acgtttgtca	ttttatttct	cctcttattt	aaattatttg	ctttctgcaa	ttgcgatttg	240
tag						243
<210> 17 <211> 237 <212> DNA		_				
<213> Star	phylococcus	aureus Bact	teriophage 4	14 AHJD		
<400> 17				•		
atgaaagttg	acgacattgt	taccttacgt	gtcaaagg.tt	atatacttca	ttacttagat	60
gatgataatg	aatacattga	ggaatttta	ccacttcacg	agtatcattt	aaccaaaaca	120
caagcaaaag	aattattacc	agacacatgt	aaactattgt	ccactacacg	cacaacgaaa	180
acaattcaag	tttattacaa	tgatttacta	caaatcgcaa	ttgcagaaag	caaataa	237
<210> 18 <211> 222 <212> DNA <213> Stag	phylococcus	aureus Bact	ceriophage 4	14 AHJD		
<400> 18						
atggaaagat	taaaattgct	tctgctggta	taccgaaaaa	cgcctttgat	acaagcgtcg	60
attttgaaac	ctttgtacgt	gaacaattct	ttgacggtgc	cattattgaa	aacaataaaa	120
gtatctataa	tgagcaaggt	acaatatcga	tatatccgtc	taaaactgaa	attgtatgtg	180
gtaatgtata	tgatgaatat	tttactgatg	aacttaatat	ga		222
<210> 19 <211> 213 <212> DNA <213> Star	phylococcus	aureus Bact	ceriophage 4	14 AHJD	·	
<400> 19 atgttaattg	gtactgtgtc	cataatcacg	tattcttcac	tatattgtcc	aataaaatct	60
tgctctttag	ctaaccaatt	aaaacgatta	cctaatgcaa	tatcgattaa	taaagtctca	120
	ggaataaata	_	_	_	- -	180
_			-		_ 9	
catttatcac	caaatgtcca	agatttgaa	taa			213

<210> <211> <212> <213>	20 207 DNA Star	phylococcus	aureus Bact	teriophage 4	44 AHJD		
<400> atgttac	20 cctq	gtttgtataa	gtattctttt	ttgaataaag	gtacaccaat	tgctttttta	60
_	_	gtaactgtgc			,	-	120
tttggct	ttga	ctgatttacc	actaattggt	ttatggtctc	cgtcatcatc	agtaggatta	180
gaactad	ctac	tcccactatc	tacttga				207
<210> <211> <212> <213>	21 189 DNA Star	phylococcus	aureus Bact	teriophage 4	44 AHJD		
<400> atgtcta	21 aaac	gattttgttt	taccatgttt	ttgctccttg	taatagttta	tgatgtcgtt	60
tacagto	gtta	aatttattcg	tcaaatgttg	cataatataa	aaagttatac	ctcacatctt	120
catcato	caat	atttgtcact	ggtctatctg	atttaccaat	ttctttatat	aaagtatcga	180
tttcttt	caa						189
<210><211><211><212><213>	22 183 DNA Star	phylococcus	aureus Bact	teriophage 4	44 AHJD		
<400>	22					•	C 0
_		aaagtcaaca	-	_			60
		catttatatc				_	120
atgtttt	tgtt	tccgcattaa	cattactgaa	gaattcttta	tattccgata	tattagcctc	180
taa							183

<211>	23 144 DNA						
		hylococcus	aureus Bact	eriophage 4	44 AHJD		
<400> 2 atgtttg		aaatgataat	acagaatatc	aataatttt	tagaaaatcc	tctcattgat	60
ttttttga	acc	ataagttatt	atttttaatt	gcttttgaaa	tacctgtaat	aatatcaacg	120
aacattaa	ata	caaataaaaa	gtag				144
<211> 3 <212> 1	24 180 DNA Stap	hylococcus	aureus Bact	ceriophage 4	44 AHJD		
	24 cac	ccccaagga	ataccaacac	tgtaactatt	acctgttttt	ccattccatt	60
ggcgcact	tgg	taaataataa	cgtgtgcctt	gccagttata	accaatccat	acgtaaccat	120
ctgataaa	aca	aacttcgtta	tatggtgtat	aaccgtttgg	ttggaaccaa	tagccattag	180
<212> I	177 DNA	hylococcus	aureus Bact	ceriophage 4	44 AHJD		
	25						
gtgtcaat	tgt	acgcctcttg	taaatcttta	tcatcaaatt	taaaattaac	attactaaaa	60
tcatttaa	aaa	ataaatcttt	ttcttgctct	tttctagctt	ctctttcttt	tttccatcta	120
tccattto	cag	acgtatgtct	aaccaatgtt	atcaacctcc	atataaagca	taaataa	177
<211> 3 <212> 1 <213> 3	_	hylococcus	aureus Bact	ceriophage 4	44 AHJD		
	26 gta	aatacaaaac	ggtattatta	tattgcgatg	agattaaagg	acattttcca	60
catcaaa	tct	caatgtttga	agatttatat	gacgctaaag	ttgtatattc	atattatgaa	120
tataacci	tat	tcactaaaaa	atacgcgtat	atcatagaat	acattaagga	gatataa	177

	<210> <211> <212>	27 168 DNA						
	<213>	Stap	hylococcus	aureus Bac	cteriophage	44 AHJD		
	<400>	27	tattaaacat	agccattgti	t ttccttttag	catttttaat	tacacttatc	60
	acgaaca			agoodogo				
	atactta	atga	cactgcatat	acgcgtgtca	a tttggtgttt	tattcactac	attgattata	120
	ttctata	atta	tctttttaat	ggttatttat	gctttatatg	gaggttga		168
		28 165 DNA Star	phylococcus	aureus Bac	cteriophage	44 AHJD		
	.400>	0.0						
	<400> atgatto	28 gtct	atatccctaa	ttttagtaca	a aaattcatat	tgttttgtat	atggtacaac	60
	gataata	attt	gtcataaaag	tagttacati	atacatgact	ttaatatatt	tatcatcagt	120
	tttgata	atag	aagaaatcac	cgttttgatt	gatgtgattt	cttaa		165
•						•		
	<210>	29	•				•	
	<211>	165						
	<212>	DNA						
	<213>	Star	hylococcus	aureus Bac	cteriophage	44 AHJD		
	<400>	20						
	<400>	29	taasaataas	2++a+2aa+	g ctttcatatt	++++ a aaaaa	tataaattaa	60
	atyyaat	Lala	tycacytcca	actiguacti	y Cilicalati	ttttgcaaaa	tetgeattae	00
	cttttct	ttg	tacgtcttgt	ggtacaaagt	ggacgatgtt	acctgcgtca	taccaagacg	120
	gttgtcd	cagc	ttgttttgat	tgtgatacta	a actttcttgc	tatga		165
	<210>	30						
	<211>	165						
	<212>	DNA						
	<213>	Stap	hylococcus	aureus Bac	cteriophage	44 AHJD		
	<400>	30						
	gtgttta	aaat	ggaacgtaaa	tacaaaacg	g tattattata	ttgcgatgag	attaaaggac	60
	attttc	caca	tcaaatctca	atgtttgaag	g atttatatga	cgctaaagtt	gtatattcat	120
	attatos	asta	taacctattc	actaaaaaa	t acqcqtatat	catad		165

	<210> <211>	31 162						
	<212> <213>	DNA Stap	phylococcus	aureus Bac	teriophage	44 AHJD		
	<400>	31						
			caccaaatga	cacgcgtata	tgcagtgtca	taagtatgat	aagtgtaatt	60
	aaaaat	gcta	aaaggaaaac	aatggctatg	tttaataggt	tattcatggt	caatcacttt	120
	cccatta	atcg	tatatgactt	tgttttgata	aataatcatt	aa		162
	<210><211><211><212><213>	32 153 DNA Star	phylococcus	aureus Bac	teriophage	44 AHJD		
	<400>	32						
			atagttcatt	gttatcatct	aaacggaata	agttaaaatg	tgaacgtaat	60
	gcaggta	atgc	catataatcc	atttaaaacg	actttagata	acataacctc	ctcatttgag	120
that that them tam that the that	tatggg	tgtt	cgttgatatc	atcagtaatg	tga			153
Tiper.	<210><211><211><212><213>	33 150 DNA Stag	phylococcus	aureus Bac	teriophage :	44 AHJD		
	<400>	33						
a first strate			ttgctaaagc	gagtagtgaa	ctaccactgt	caccactact	accactgtca	60
j	gacgaat	tcac	taggtgatcc	acctttaccg	tctaatttac	caccccaagc	tagaatagta	120
	ttcgca	ccgt	ctaaaaatgg	attaccatag	•			150
	<210><211><211><212><213>	34 150 DNA Stap	phylococcus	aureus Bac	teriophage	44 AHJD	-	
	<400>	34						
			ttaaccacct	ctaccaaatt	tgtaaaaaac	attttttatc	aaattcattt	60
	aaaatt	ttct	ttcttaaatc	gtacgcttta	tcaatattat	caattaaata	ctgcttagtg	120
	aattgt	gtac	cttttgcatt	acctttttga				150

<210> 35 <211> 14	7					
<212> DNZ <213> Sta	A aphylococcus	aureus Baci	teriophage :	44 AHJD		
(210)	ap, 100000as	darous sus	301 2 0pa90			
<400> 35						
atgatgatt	c taataataaa	aacgctaaaa	agcataaata	cgctttatat	aatttacaag	60
ctaaaaata	a taattcttca	atgtataaat	atattaaaga	aatcgatact	ttatataaag	120
aaattggta	a atcagataga	ccagtga				147
			•			
<210> 36						
<211> 14						
<212> DN	A					
<213> St	aphylococcus	aureus Bact	teriophage	44 AHJD		
<400> 36						
	t tgacgaataa	atttaacact	gtaaacgaca	tcataaacta	ttacaaggag	60
caaaaacat	g gtaaaacaaa	atcqtttaga	catggtaaga	gattatcaaa	atgctgtcaa	120
tcatgtcag	a aaaaaaatcc	cagataa				147
<210> 37	_		•			
<211> 14						
<212> DNZ				4.4 711.TD		
<213> Sta	aphylococcus	aureus Baci	teriophage	44 AHJD		
<400> 37						
gtgtataca	a taccacacgt	gatggtgcaa	catatggtgg	tacattatag	tttgcaacta	60
aaaacgaac	c atcttcaaaa	actgctacaa	caacacctgt	gtgaccaata	ccatatgcag	120
ttacttata	a gtatggtggt	ttactac				147
cegeeegea	a geaeggegge	ccaccag				117
<210> 38	,					
<210> 38 <211> 14	1					
<211> 14						
	aphylococcus	aureus Bact	teriophage	44 AHJD		
	apily rococcus	dareas bac	corrobinage			
<400> 38						
atgtcgata	t ctaacgtaaa	taactctttt	tcaatttcaa	aatcatcata	ttgtttgtca	60
aactcaata	t acacatcacc	catatttatt	tttactatac	attttttatt	agatgaagta	120
aattttca	a atttatcatt	ataa				144

	39 144 DNA						
<213>	Stap	ohylococcus	aureus Bac	teriophage -	44 AHJD		
<400> gtgtaco	39 cttt	tgcattacct	ttttgatttt	gattacgttt	tgcgttttga	ttactttcgt	60
tactcga	attt	attcacagtt	ttaccgttat	caatcgtatt	attatcagcg	aatcgtaacg	120
ttgtatt	tatc	aacatcaatg	ttaa				144
<211>	40 141 DNA Star	phylococcus	aureus Bac	teriophage ·	44 AHJD		
<400>	40						
gtgctgt	att	tacttatgat	gtatctaaac	ttaaagagtt	tactggcaac	gttgaagaaa	60
ttaaaco	caaa	atcagattta	tatgcgttta	ttttggatat	taattcaatt	aaatataaac	120
gttacad	caaa	aggtatgtta	a				.141
<210>	41						
<211> <212>	138 DNA						
<213>	Stap	hylococcus	aureus Bac	teriophage ·	44 AHJD		
<400> gtggtaa	41 actg	gacatatgca	cagttaccag	aaaaatataa	aaaagcaatt	ggtgtacctt	60
tattcaa	aaaa	agaatactta	tacaaaccag	gtaacatatt	tcctcaaacg	ggtaatgcag	120
gacaato	gtac	agaattaa					138
<210> <211> <212> <213>	42 138 DNA Stap	phylococcus	aureus Bac	teriophage ·	44 AHJD		
<400>	42						
atgtcgt	caa	ctttcattat	tatatcactc	ctttctaaaa	aacgtaaacg	ttatacgttt	60
cataaaa	atcc	tttatgcata	ttccattgtt	ctattgggtc	atcaccagca	atataagaca	120
atattga	attc	tggtttag					138

	43 138 DNA					
	Staphylococcus	aureus Bacte	eriophage 4	4 AHJD		
<400>	43 acc gtcgtctttt	gttaatttat a	agttttgtga	acctcttgcg	cataatactt	60
caaagtg	ttc atactcacca	agttggaaga a	accatataa	attatggaaa	cgttttccac	120
caccgcc	gtt tgtcatag	•				138
<212>	44 138 DNA Staphylococcus	aureus Bacte	eriophage 4	4 AHJD'		
<400>	44 ttg taacagtttt	ggaagagat g	ratastatss	ccagattttc	atttcaccat	60
		-				
tggattg	gacg ttctaatccg	attgttgtac o	catgaccacc	ctgtacaata	cgcatgcttg	120
aaattaa	igtc accactag					138
<212>	45 135 DNA Staphylococcus	aurous Bacte	arionhage 1	4 AHID		
		aureus bacce	silophage 4	4 ANOD		
<400>	45 cta tttacgtgat	gatatgtttt a	ataaagaaaa	catggaacgt	tattactaca	60
_	caa tttacatttt					120
gatattt	ata tttag					135
J	.					
<210> <211> <212>	46 129 DNA					·
<213>	Staphylococcus	aureus Bacte	eriophage 4	4 AHJD		
<400> atggcac	46 cgt caaagaattg	ttcacgtaca a	aaggtttcaa	aatcgacgct	tgtatcaaag	60
acattt	tcg gtataccage	agaagcaatt t	taatctttc	cattcacttc	atatgcatat	120
		, <u>j j</u>			. 5	129
ttcttat	:ga					129

	֡
1	
1	
1	
¥	
Ü	
į.	
ļ.£	

	47 129						
	DNA Stap	phylococcus	aureus Bac	teriophage 4	14 AHJD		
<400>							60
atgatta	atcc	atttaagtta	tcatatcaag	acggtattaa	tttcccacgt	gataacttta	60
aagagco	ctga	gggtatttgc	atttatacaa	atccaaaaac	aaaacgtaaa	tcgttattac	120
ttgctat	ga						129
	48 129						
	DNA	1 1	D		4.4 311.75		
<213>	Star	phylococcus	aureus Bac	teriophage 4	44 AHJD		
<400>	48						
atgaatg	gtat	gtaagttgtt	caggtgtgag	ttttgcaaaa	catttcacag	catagtcata	60
ggcttca	acta	tcattcatat	cattatcttt	atcaaaaatc	gtataattaa	aatctgtttt	120
aagttgt	ga						129
<210>	49						
	129						
	DNA		_				
<213>	Stap	phylococcus	aureus Bac	teriophage 4	14 AHJD		
<400>	49						
atgagga	acgt	tgttgacatt	atcaatgctg	gagaagttca	attcacaatt	tatgaatatg	60
aaaacaa	aaaa	aggtcaaaaa	ggttactcaa	tcaattttgg	tcaagtatca	ttttaataca	120
atttcat	ag						129
<210>	50						
<211>	126						
<212>	DNA						
<213>	Stap	phylococcus	aureus Bac	teriophage 4	14 AHJD		
<400>	50						
atgaggg	gggc	aggtgttgac	tttgatggtg	catatggatt	tcaatgtatg	gacttatcag	60
ttgctta	atgt	gtattacatt	actgacggta	aagttcgcat	gtggggtaat	gctaaagacg	120
cgataa							126

Ø
إلى
ΠJ
الم
Ü
T
IJ
E

	<210> <211> <212>	51 126 DNA							
	<213>		hylococcus	aureus	Bact	eriophage (44 AHJD		
	<400>	51		•					
	gtgtgtt	cacg	tttttcattc	acgtaat	cgt	ttcgtcgcat	ttctaaaaaa	atgtttttgt	60
	aaagtct	tga	tgtattcatt	ttatgct	ttt	gtaataaatt	gtatatattt	aaattggata	120
	atatag								126
	3								
	<210>	52							
	<211>	123						•	
	<212>	DNA							
	<213>	Stap	hylococcus	aureus	Bact	eriophage 4	44 AHJD		
	<400>	52							
	atgataa	acaa	tgaactatac	aatatca	tta	acggttacaa	aaacactgaa	cgtaatatat	60
7	+ - + - - + -			.		****		++====+==+	120
	tattete	ctac	attigicaca	ccacgii	Cat	tgtataactt	attggttcct	LLCCAALACL	120
term Land Tout (Ern Min thin this Mill Ma	taa								123
ý *									
.) 기	<210>	53							
ni H	<211>	123							
zr	<212>	DNA							
17. 17.	<213>	Stap	hylococcus	aureus	Bact	eriophage 4	44 AHJD		
] ≟	<400>	53							
			taatgttaat	tttaaat	ttg	atgataaaga	tttacaagag	gcgtacattg	60
ž.	acacato	ggaa	acattttgca	catctgc	cct	attttcctaa	agaaagaaac	gtatcatatg	120
	taa								123
	<210>	54							
	<211>	120							
	<212>	DNA							
	<213>		hylococcus	aureus	Bact	eriophage	44 AHJD		
			3			1 3			
	<400>	54							
	atgtggt	tta	ttcatcaagt	gaagttg	aaa	aatacttaca	atcacaaggc	ttcacagaac	60
	acaatga	aaga	tacaacaagt	aacacto	atg	aaacatcgaa	tcaaaatgct	acatctttag	120

<210>	55						
<211>	120						
<212>	DNA						
<213>	Stap	phylococcus	aureus Ba	cteriophage	44 AHJD		
<400>	55						
atgact	ggaa	tggaaatacg	atgttactc	g acgctggtaa	gatttcacaa	aaaactggtg	60
ttaagt	tacg	tacaaaatca	attattggt	t atcataatga	agttcgagta	tatccagtag	120
			_		44		
<213>	Stap	phylococcus	aureus Ba	cteriophage	44 AHJD		
4400>	5 C						
						.	60
atgtgt	ctgg	taataattct	tttgcttgt	g titiggitaa	atgatactcg	tgaagtggta	60
				<u> </u>			117
aaaatt	cctc	aatgtattca	ttatcatca	t ctaagtaatg	aagtatataa	cctttga	117
<210×	E 7						
		obylogoggus	aurous Ba	ctoriophago	44 VA TD		
\213/	Stap	onyrococcus	aureus ba	ccerrophage	44 ANOD		
<4005	57						
		cattacaddt	aaccaaato	a aattattaa	anacnencea	gaagaatta	60
gegage	acca	caccacagge	aaccaaacy	g dattattag	agacgcgcca	gaagaaacca	00
aaaaag	taaa	tacataatta	catatatca	a gtggtaacgc	agtcggtgaa	gtaa	114
aaaaag	-999	cycacyycca	cgcgcgcca	a geggeaaege	ageeggegaa	gcaa	
<210>	58						
		ohvlococcus	aureus Ba	cteriophage	44 AHJD		
	1						
<400>	58						
atgtac	cacc	atatqttqca	ccatcacqt	g tggtattgta	tacactcatt	aatggcgtac	60
,		, ,	,	<i>3</i>		33 3	
caaata	atgc	tggtgataat	attgtattc	t ttagtggtat	tgcttaatta	a	111
	_		-		-		
<210>	59						
<211>	111						
<212>	DNA						
<213>	Stap	phylococcus	aureus Ba	cteriophage	44 AHJD		
<400>	59						
atgcat	attt	cttatgattc	agtacaaac	a tcttatctat	ctgttcgttt	tcaatatccc	60
	<211> <212> <213> <400> atgact ttaagt <210> <211> <212> <213> <400> atgttt aaaatt <210> <211> <212> <213> <400> gtgagt aaaaag <210> <211> <212> <213> <400> gtgagt aaaaag <210> <211> <212> <213> <400> atgtt <210> <211> <212> <213> <400> atgtac caaata <210> <211> <212> <213> <400> atgtac caaata <210> <211> <212> <213> <400> atgtac caaata <210> <211> <212> <213> <210> <211> <212> <213> <210> <210> <211> <212> <213> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <	<pre><211> 120 <212> DNA <213> Staj <400> 55 atgactggaa ttaagttacg <210> 56 <211> 117 <212> DNA <213> Staj <400> 56 atgtgtctgg aaaattcctc <210> 57 <211> 114 <212> DNA <213> Staj <400> 57 gtgagtatta aaaagtggg <210> 58 <211> 111 <212> DNA <213> Staj <400> 57 gtgagtatta aaaagtggg <210> 58 <211> 111 <212> DNA <213> Staj <400> 58 atgtaccacc caaataatgc <210> 58 <211> 111 <212> DNA <213> Staj <400> 58 atgtaccacc caaataatgc</pre>	<pre><211> 120 <212> DNA <213> Staphylococcus <400> 55 atgactggaa tggaaatacg ttaagttacg tacaaaatca <210> 56 <211> 117 <212> DNA <213> Staphylococcus <400> 56 atgtgtctgg taataattct aaaattcctc aatgtattca <210> 57 <211> 114 <212> DNA <213> Staphylococcus <400> 57 gtgagtatta cattacaggt aaaagtggg tgcatggtta <210> 57 gtgagtatta cattacaggt aaaaagtggg tgcatggtta <210> 58 <211> 111 <212> DNA <213> Staphylococcus <400> 58 atgtaccacc atatgttgca caaataatgc tggtgataat <210> 58 atgtaccacc atatgttgca caaataatgc tggtgataat <210> 58 <211> 111 <212> DNA <213> Staphylococcus <400> 58 atgtaccacc atatgttgca caaataatgc tggtgataat <210> 59 <211> 111 <212> DNA <213> Staphylococcus <400> 59</pre>	<pre><211> 120 <212> DNA <213> Staphylococcus aureus Ba <400> 55 atgactggaa tggaaatacg atgttactc ttaagttacg tacaaaatca attattggt <210> 56 <211> 117 <212> DNA <213> Staphylococcus aureus Ba <400> 56 atgtgtctgg taataattct tttgcttgt aaaattcctc aatgtattca ttatcatca <210> 57 <211> 114 <212> DNA <213> Staphylococcus aureus Ba <400> 57 gtgagtatta cattacaggt aaccaaatg aaaaagtggg tgcatggtta cgtgtgtca <210> 57 gtgagtatta cattacaggt accaaatg aaaaagtggg tgcatggtta cgtgtgtca <210> 58 <211> 111 <212> DNA <213> Staphylococcus aureus Ba <400> 58 atgtaccacc atatgttgca ccatcacgt caaataatgc tggtgataat attgtattc <210> 58 atgtaccacc atatgttgca ccatcacgt caaataatgc tggtgataat attgtattc <210> 59 <211> 111 <212> DNA <213> Staphylococcus aureus Ba <400> 59</pre>	<pre><211> 120 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 55 atgactggaa tggaaatacg atgttactcg acgctggtaa ttaagttacg tacaaaatca attattggtt atcataatga <210> 56 <211> 117 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 56 atgtgtctgg taataattct tttgcttgtg ttttggttaa aaaattcctc aatgtattca ttatcatcat ctaagtaatg <210> 57 <211> 114 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 57 gtgagtatta cattacaggt aaccaaatgg aattatttag aaaaagtggg tgcatggtta cgtgtgtcaa gtggtaacgc <210> 57 gtgagtatta cattacaggt cgtgtcaa gtggtaacgc <210> 58 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 58 atgtaccacc atatgttgca ccatcacgtg tggtattgta caaataatgc tggtgataat attgtattct ttagtggtat <210> 58 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 58 atgtaccacc atatgttgca ccatcacgtg tggtattgta caaataatgc tggtgataat attgtattct ttagtggtat <210> 59 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage <400> 59 </pre>	<pre><211> 120 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 55 atgactggaa tggaaatacg atgttactcg acgctggtaa gatttcacaa ttaagttacg tacaaaatca attattggtt atcataatga agttcgagta </pre> <pre><210> 56 <211> 117 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 56 atgtgtctgg taataattct tttgcttgtg ttttggttaa atgatactcg aaaattcctc aatgtattca ttatcatcat ctaagtaatg aagtatataa </pre> <pre><210> 57 <211> 114 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 57 gtgagtatta cattacaggt aaccaaatgg aattatttag agacgcgcca aaaaagtggg tgcatggtta cgtgtgtcaa gtggtaacgc agtcggtgaa </pre> <pre><210> 58 <211> 111 <212> DNA </pre> <pre><210> 58 atgtaccacc atatgttgca ccatcacgtg tggtattgta tacactcatt caaataatgc tggtgataat attgtattct ttagtggtat tgcttaatta </pre> <pre><400> 58 atgtaccacc atatgttgca ccatcacgtg tggtattgta tacactcatt caaataatgc tggtgataat attgtattct ttagtggtat tgcttaatta </pre> <pre><210> 59 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD</pre> <400> 59	<pre><211> 120 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 55 atgactggaa tggaaatacg atgttactcg acgctggtaa gatttcacaa aaaactggtg ttaagttacg tacaaaatca attattggtt atcataatga agttcgagta tatccagtag <210> 56 <211> 117 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 56 atgtgtctgg taataattct tttgcttgtg ttttggttaa atgatactcg tgaagtggta aaaattcctc aatgtattca ttatcatcat ctaagtaatg aagtatataa cctttga <210> 57 <211> 114 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 57 gtgagtatta cattacaggt aaccaaatgg aattatttag agacgcgca gaagaaatta aaaaagtggg tgcatggtta cgtgtgtcaa gtggtaacgc agtcggtgaa gtaa <210> 58 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 58 atgtaccacc atatgttgca ccatcacgtg tggtattgta tacactcatt aatgcgtac caaataatgc tggtgataat attgtattct ttagtggtat tgcttaatta a </pre> <pre><210> 59 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD</pre>

T
ij
ij
IJ
E
Ü
-
ļu.Ēs

	atttaco	ctaa	ggctatcggg	tcgaataaac	tggggttcaa	taagggttta	a	111
	<210>	60						
	<211>	111						
	<212>	DNA						
	<213>		phylococcus	aureus Bact	eriophage 4	44 AHJD		
	<400>	60						
	atggatt	ttg	taacattgga	ttacctgaac	cgtcattatg	ccaaaatctt	acaccagatt	60
	ctaaaat	tgc	ttttaattgt	tccattaaca	tggggtcgat	gtcacgtata	g	111
	<210>	61						
	<211>	111						
		DNA				•		
	<213>		phylococcus	aureus Bact	eriophage 4	14 AHJD		
	<400>	61						
	atgtaco	catt	ttcatttcta	taatatgtgc	cgtattggtt	tcgtttccat	tttccaaatg	60
	tatttac	cttt	tgatgtttct	aatgctttgc	tattactacc	tgaaaattta	g	111
j	<210>	62						
Ō	<211>	108						
Ü	<212>	DNA						
	<213>		phylococcus	aureus Bact	eriophage 4	14 AHJD		
) j	<400>	62						
rk:			gtgtcttgat	aaaatatctt	ttacgtttgt	cattttattt	ctcctcttat	60
	ttaaatt	tatt	tgctttctgc	aattgcgatt	tgtagtaaat	cattgtaa		108
	<210>	63				•	•	
	<211>	105						
	<212>	DNA						
	<213>	Stap	ohylococcus	aureus Bact	eriophage 4	14 AHJD		
	<400>	63						
	gtggtat	tcg	caacgcagtt	aaccaatcta	ttaatattga	taaagaaaca	aatcacatgt	60
	actctad	caca	atccgattct	caaaaacctg	aaggtttttg	gataa		105

<210> <211> <212>	64 105 DNA		
<213>	Staphylococcus	aureus Bacteriophage 44 AHJD	
<400> atgcgto	64 cttg tattttttt	aataattctt gcatggcttg ttttgctaaa gcgagtagtg	60
aactac	cact gtcaccacta	ctaccactgt cagacgaatc actag	105
<210> <211> <212> <213>	102 DNA	aureus Bacteriophage 44 AHJD	
<400>	65		
atgacga	agtc aatcaatcaa	cttgtgtccg aaatatataa cggtgcacca tttgttaaaa	60
tgtcaco	ctat gtttaatgca	gatgacgata tcattgattt aa	102
<210> <211> <212> <213>		aureus Bacteriophage 44 AHJD	
<400>	66		
gtggtg	gaaa acgtttccat	aatttatatg gtttcttcca acttggtgag tatgaacact	60
ttgaag	catt acgcgcaaga	ggttcacaaa actataaatt aa	102
<210><211><211><212><213>	67 102 DNA Staphylococcus	aureus Bacteriophage 44 AHJD	
<400>	67		
atgata	tct ttatattgaa	agtgacatcg gttcattttc acttaacgac ttatttccag	60
ttgaac	gttc agtacataac	aaatctgatt tgcatatatt aa	102
<210><211><211><212><213>		aureus Bacteriophage 44 AHJD	
<400>	68		
	gaat ttgatgaaat	cgtaaaacca gacgacaaag aagaaacttc agaatcaact	60

gaagaaa	aatt	tagaatcaac	tgaagaaact	tcagaatcaa	ctgaagaatc	aactgaagaa	120
gaagaa		cagaaccaac	ogaagaaaoo				
tcaacto	gaag	aatcaactga	agataaaaca	gtagaaacaa	tcgaagaaga	aaatgaaaac	180
aaattag	gaac	ctactacaac	agatgaagat	agttcgaaat	ttgaccctgt	tgtattagaa	240
caacgta	attg	cttcattaga	acaacaagtg	actacttttt	tatcttcaca	aatgcaacaa	300
ccacaa	caag	tacaacaaac	acaatcagat	gtaacagaat	caaacaaaga	agataacgac	360
tattcagatg		aagaactagt	tgataagtta	gatttagatt	ag		402
<210> <211> <212> <213>	69 303 DNA Star	phylococcus	aureus Baci	teriophage 4	14 AHJD		
<400> atggtta	69 aatg	ttgataatgc	accagaagaa	aaaggacaag	cctatactga	aatgttgcaa	60
ctattca	aata	aactgattca	atggaatcca	gcttatacat	ttgacaatgc	aattaactta	120
ttatcgg	gctt	gccaacaact	attattaaac	tataatagtt	ctgttgttca	attcttaaat	180
gatgaad	ctaa	acaacgaaac	taaaccagaa	tcaatattgt	cttatattgc	tggtgatgac	240
ccaatag	gaac	aatggaatat	gcataaagga	ttttatgaaa	cgtataacgt	ttacgttttt	300
tag							303
<210> <211> <212> <213>	70 198 DNA Star	phylococcus	aureus Bact	teriophage 4	14 AHJD		
<400>	70						
atggaaa	aatg	aaacaaaaaa	cattgagttg	aagcatgttt	ttcgttttaa	gaatggaagt	60
ttatgta	atag	cgttatttga	tagaacagaa	aatgaaattt	cattttatga	tgttgacatt	120
gatgaaa	attg	aagatttaaa	tcataattct	gttttacgcg	taatttcaac	tttattagga	180

agtgataata atggttaa

<211>	183						
<212> <213>		phylococcus	aureus Bac	teriophage 4	44 AHJD		
<400>	71						
atgtate	gagg	gaaacaacat	gcgttctatg	atgggtacat	catatgaaga	ttcaagatta	60
aataaa	cgaa	cagaattaaa	tgaaaacatg	tcaattgata	caaataaaag	tgaagatagt	120
tatggt	gtac	aaattcattc	actttcaaaa	caatcattta	caggtgacgt	tgaggaggaa	180
taa							183
<21.0×	70						
<213>		hylococcus	aureus Bact	teriophage 4	44 AHJD		
<400>	72						
	acat	gtcaattgat	acaaataaaa	gtgaagatag	ttatggtgta	caaattcatt	60
cacttt	caaa	acaatcattt	acaggtgacg	ttgaggagga	ataataaatt	atggcacaac	120
aatcta	caaa	aaatgaaact	gcacttttag				150
						•	
<210>	73						
<213>	Stap	phylococcus	aureus Bact	teriophage 4	14 AHJD		
<400>	73						
atgatto	gttt	tgaaagtgaa	tgaatttgta	caccataact	atcttcactt	ttatttgtat	60
caattga	acat	gttttcattt	aattctgttc	gtttatttaa	tcttgaatct	tcatatgatg	120
tacccat	tcat	ag					132
<210S	74						
				•			
<213>		hylococcus	aureus Bact	teriophage 4	14 AHJD		
<400>	74						
		ttaattctgt	tcgtttattt	aatcttgaat	cttcatatga	tgtacccatc	60
atagaad	cgca	tgttgtttcc	ctcatacatg	tttaaattcc	tcctaatcta	a	111
	<211> <212> <213> <400> atgtate aataaa tatggte taa <210> <211> <212> <213> <400> atgaaa cacttte aatctae <210> <211> <212> <213> <400> atgate tacccae <210> <211> <212> <213> <400> atgate tacccae <210> <211> <212> <213> <400> atgate tacccae	<212> DNA <213> Stap <400> 71 atgtatgagg aataaacgaa tatggtgtac taa <210> 72 <211> 150 <212> DNA <213> Stap <400> 72 atgaaaacat cactttcaaa aatctacaaa <210> 73 <211> 132 <212> DNA <213> Stap <400> 73 atgattgtt caattgacat tacccatcat <210> 74 <211> 111 <212> DNA <213> Stap <400> 73 atgattgttt caattgacat tacccatcat <210> 74 <211> 111 <212> DNA <213> Stap <400> 74 atgttttcat	<pre><211> 183 <212> DNA <213> Staphylococcus <400> 71 atgtatgagg gaaacaacat aataaacgaa cagaattaaa tatggtgtac aaattcattc taa <210> 72 <211> 150 <212> DNA <213> Staphylococcus <400> 72 atgaaaacat gtcaattgat cactttcaaa acaatcattt aatctacaaa acaatcattt aatctacaaa aaatgaaact <210> 73 <211> 132 <212> DNA <213> Staphylococcus <400> 73 atgattgtt tgaaagtgaa caattgacat gtttcattt tacccatcat ag <210> 74 <211> 111 <212> DNA <213> Staphylococcus <400> 73 atgattgttt tgaaagtgaa caattgacat gttttcattt tacccatcat ag <210> 74 <211> 111 <212> DNA <213> Staphylococcus <400> 74 atgttttcat ttaattctgt</pre>	<pre><211> 183 <212> DNA <213> Staphylococcus aureus Bace <400> 71 atgtatgagg gaaacaacat gcgttctatg aataaacgaa cagaattaaa tgaaaacatg tatggtgtac aaattcattc actttcaaaa taa <210> 72 <211> 150 <212> DNA <213> Staphylococcus aureus Bace <400> 72 atgaaaacat gtcaattgat acaaataaaa cactttcaaa acaatcatt acaggtgacg aatctacaaa aaatgaaact gcacttttag <210> 73 <211> 132 <212> DNA <213> Staphylococcus aureus Bace <400> 73 atgattgtt tgaaagtgaa tgaattgta caattgacat gtttcatt aattctgttc tacccatcat ag <210> 74 <211> 111 <212> DNA <213> Staphylococcus aureus Bace <400> 74 <211> 111 <212> DNA <213> Staphylococcus aureus Bace <400> 74 atgttttcat ttaattctgt tcgtttattt</pre>	<pre><211> 183 <212> DNA <213> Staphylococcus aureus Bacteriophage 4 <400> 71 atgtatgagg gaaacaacat gcgttctatg atgggtacat aataaacgaa cagaattaaa tgaaaacatg tcaattgata tatggtgtac aaattcattc acttcaaaa caatcatta taa <210> 72 <211> 150 <212> DNA <213> Staphylococcus aureus Bacteriophage 4 <400> 72 atgaaaacat gtcaattgat acaaataaaa gtgaagatag cacttcaaa acaatcatt acaggtgacg ttgaggagga aatctacaaa aaatgaaact gcactttag <210> 73 <211> 132 <211> 132 <212> DNA <213> Staphylococcus aureus Bacteriophage 4 <400> 73 atgattgtt tgaaagtgaa tgaattgta caccataact caattgacat gtttcatt aattctgttc gtttattaa tacccatcat ag <210> 74 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 4 <400> 74 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 4 <400> 74 atgtttcat ttaattctgt tcgtttattt aatcttgatt</pre>	<pre><211> 183 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 71 atgtatgagg gaaacaacat gcgttctatg atgggtacat catatgaaga aataaacgaa cagaattaaa tgaaaacatg tcaattgata caaataaaag tatggtgtac aaattcattc actttcaaaa caatcattta caggtgacgt taa </pre> <pre><210> 72 <211> 150 <212> DNA </pre> <pre><213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 72 atgaaaacat gtcaattgat acaaataaaa gtgaagatag ttatggtgta cactttcaaa acaatcattt acaggtgacg ttgaggagga ataataaatt aatctacaaa aaatgaaact gcacttttag </pre> <pre><210> 73 <211> 132 <212> DNA </pre> <pre><210> 73 atgattgttt tgaaagtgaa tgaattgta caccataact atcttcactt caattgacat gtttcattt aattctgttc gtttatttaa tcttgaatct tacccatcat ag </pre> <pre><400> 74 <211> 111 <212> DNA </pre> <pre><210> 74 <211> 111 <212> DNA</pre> <pre><210> 74 <211</pre>	<pre><211> 183 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 71 atgtatgagg gaaacaacat gcgttctatg atgggtacat catatgaaga ttcaagatta aataaacgaa cagaattaaa tgaaaacatg tcaattgata caaataaaag tgaagatagt tatggtgtac aaattcattc actttcaaaa caatcattta caggtgacgt tgaggaggaa taa <210> 72 <211> 150 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <pre><400> 72 atgaaaacat gtcaattgat acaaataaaa gtgaagatag ttatggtgta caaattcatt cactttcaaa acaatcattt acaggtgacg ttgaggagga ataataaatt atggcacaac aatctacaaa aaatgaaact gcacttttag <210> 73 <211> 132 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD </pre> <400> 73 atgattgtt tgaaagtgaa tgaattgta caccataact atcttcactt ttatttgtat caattgacat gtttcattt aattctgtc gtttatttaa tcttgaatct tcatatgatg tacccatcat ag <210> 74 <211> 111 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD

```
<210> 75
<211> 761
<212> PRT
```

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 75

Met Gly Leu Leu Glu Cys Met Gln Tyr His Lys His Glu Arg Arg Met 1 5 10 15

Ile Leu Tyr Trp Asp Ile Glu Thr Leu Ala Tyr Asn Lys Val Asn Gly 20 25 30

Arg Lys Lys Pro Thr Lys Tyr Lys Asn Val Thr Tyr Ser Val Ala Ile 35 40 45

Gly Trp Phe Asn Gly Tyr Glu Ile Asp Val Glu Val Phe Pro Ser Phe 50 55 60

Glu Ser Phe Tyr Asp Ala Phe Tyr Thr Tyr Val Lys Arg Arg Asp Thr 65 70 75 80

Ile Thr Lys Ser Lys Thr Asp Ile Ile Met Ile Ala His Asn Cys Asn 85 90 95

Lys Tyr Asp Asn His Phe Leu Leu Lys Asp Thr Met Arg Tyr Phe Asp 100 105 110

Asn Ile Thr Arg Glu Asn Ile Tyr Leu Lys Ser Ala Glu Glu Asn Glu 115 120 125

His Thr Leu Lys Met Lys Glu Ala Thr Ile Leu Ala Lys Asn Gln Asn 130 135 140

Val Ile Leu Glu Lys Arg Val Lys Ser Ser Ile Asn Leu Asp Leu Thr 145 150 155 160

Met Phe Leu Asn Gly Phe Lys Phe Asn Ile Ile Asp Asn Phe Met Lys
165 170 175

Thr Asn Thr Ser Ile Ala Thr Leu Gly Lys Lys Leu Leu Asp Gly Gly 180 185 190

Tyr Leu Thr Glu Ser Gln Leu Lys Thr Asp Phe Asn Tyr Thr Ile Phe 195 200 205

Asp Lys Asp Asn Asp Met Asn Asp Ser Glu Ala Tyr Asp Tyr Ala Val 210 215 220

Lys Cys Phe Ala Lys Leu Thr Pro Glu Gln Leu Thr Tyr Ile His Asn 225 230 235 240

Asp Val Ile Ile Leu Gly Met Cys His Ile His Tyr Ser Asp Ile Phe 250 Pro Asn Phe Asp Tyr Asn Lys Leu Thr Phe Ser Leu Asn Ile Met Glu 260 265 270 Ser Tyr Leu Asn Asn Glu Met Thr Arg Phe Gln Leu Leu Asn Gln Tyr 280 Gln Asp Ile Lys Ile Ser Tyr Thr His Tyr His Phe His Asp Met Asn 295 Phe Tyr Asp Tyr Ile Lys Ser Phe Tyr Arg Gly Gly Leu Asn Met Tyr 315 Asn Thr Lys Tyr Ile Asn Lys Leu Ile Asp Glu Pro Cys Phe Ser Ile 325 Asp Ile Asn Ser Ser Tyr Pro Tyr Val Met Tyr His Glu Lys Ile Pro 340 345 350 Thr Trp Leu Tyr Phe Tyr Glu His Tyr Ser Glu Pro Thr Leu Ile Pro 360 365 Thr Phe Leu Asp Asp Asp Asn Tyr Phe Ser Leu Tyr Lys Ile Asp Lys 375 380 Asp Val Phe Asn Asp Asp Leu Leu Ile Lys Ile Lys Ser Arg Val Leu 385 390 395 400 Arg Gln Met Ile Val Lys Tyr Tyr Asn Asn Asp Asn Asp Tyr Val Asn 405 410 Ile Asn Thr Asn Thr Leu Arg Met Ile Gln Asp Ile Thr Gly Ile Asp 420 425 Cys Met His Ile Arg Val Asn Ser Phe Val Ile Tyr Glu Cys Glu Tyr 440 Phe His Ala Arg Asp Ile Ile Phe Gln Asn Tyr Phe Ile Lys Thr Gln 450 455 460 Gly Lys Leu Lys Asn Lys Ile Asn Met Thr Ser Pro Tyr Asp Tyr His 465 470 480 Ile Thr Asp Asp Ile Asn Glu His Pro Tyr Ser Asn Glu Glu Val Met 485 490 Leu Ser Lys Val Val Leu Asn Gly Leu Tyr Gly Ile Pro Ala Leu Arg 505 500

Ser His Phe Asn Leu Phe Arg Leu Asp Asp Asn Asn Glu Leu Tyr Asn Ile Ile Asn Gly Tyr Lys Asn Thr Glu Arg Asn Ile Leu Phe Ser Thr Phe Val Thr Ser Arg Ser Leu Tyr Asn Leu Leu Val Pro Phe Gln Tyr Leu Thr Glu Ser Glu Ile Asp Asp Asn Phe Ile Tyr Cys Asp Thr Asp Ser Leu Tyr Met Lys Ser Val Val Lys Pro Leu Leu Asn Pro Ser Leu Phe Asp Pro Ile Ala Leu Gly Lys Trp Asp Ile Glu Asn Glu Gln Ile Asp Lys Met Phe Val Leu Asn His Lys Lys Tyr Ala Tyr Glu Val Asn Gly Lys Ile Lys Ile Ala Ser Ala Gly Ile Pro Lys Asn Ala Phe Asp Thr Ser Val Asp Phe Glu Thr Phe Val Arg Glu Gln Phe Phe Asp Gly Ala Ile Ile Glu Asn Asn Lys Ser Ile Tyr Asn Glu Gln Gly Thr Ile Ser Ile Tyr Pro Ser Lys Thr Glu Ile Val Cys Gly Asn Val Tyr Asp Glu Tyr Phe Thr Asp Glu Leu Asn Met Lys Arg Glu Phe Ile Leu Lys Asp Ala Arg Glu Asn Phe Asp His Ser Gln Phe Asp Asp Ile Leu Tyr Ile Glu Ser Asp Ile Gly Ser Phe Ser Leu Asn Asp Leu Phe Pro Val Glu Arg Ser Val His Asn Lys Ser Asp Leu His Ile Leu Lys Arg Glu His Asp Glu Ile Lys Lys Gly Asn Cys

```
<210>
       76
<211>
       647
<212>
       PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
Met Ala Tyr Asn Glu Asn Asp Phe Lys Tyr Phe Asp Asp Ile Arg Pro
                                     10
Phe Leu Asp Glu Ile Tyr Lys Thr Arg Glu Arg Tyr Thr Pro Phe Tyr
            20
                                 25
Asp Asp Arg Ala Asp Tyr Asn Thr Asn Ser Lys Ser Tyr Tyr Asp Tyr
        35
Ile Ser Arg Leu Ser Lys Leu Ile Glu Val Leu Ala Arg Arg Ile Trp
                        55
Asp Tyr Asp Asn Glu Leu Lys Lys Arg Phe Lys Asn Trp Asp Asp Leu
                    70
                                         75
Met Lys Ala Phe Pro Glu Gln Ala Lys Asp Leu Phe Arg Gly Trp Leu
                                                         95
                85
Asn Asp Gly Thr Ile Asp Ser Ile Ile His Asp Glu Phe Lys Lys Tyr
            100
                                 105
                                                     110
Ser Ala Gly Leu Thr Ser Ala Phe Ala Leu Phe Lys Val Thr Glu Met
        115
                            120
                                                 125
Lys Gln Met Asn Asp Phe Lys Ser Glu Val Lys Asp Leu Ile Lys Asp
                        135
                                             140
Ile Asp Arg Phe Val Asn Gly Phe Glú Leu Asn Glu Leu Glu Pro Lys
                                         155
                    150
Phe Val Met Gly Phe Gly Gly Ile Arg Asn Ala Val Asn Gln Ser Ile
                165
                                                         175
Asn Ile Asp Lys Glu Thr Asn His Met Tyr Ser Thr Gln Ser Asp Ser
            180
                                 185
Gln Lys Pro Glu Gly Phe Trp Ile Asn Lys Leu Thr Pro Ser Gly Asp
                            200
Leu Ile Ser Ser Met Arg Ile Val Gln Gly Gly His Gly Thr Thr Ile
                        215
                                             220
```

Gly Leu Glu Arg Gln Ser Asn Gly Glu Met Lys Ile Trp Leu His His

230

Asp Gly Val Ala Lys Leu Leu Gln Val Ala Tyr Lys Asp Asn Tyr Val Leu Asp Leu Glu Glu Ala Lys Gly Leu Thr Asp Tyr Thr Pro Gln Ser Leu Leu Asn Lys His Thr Phe Thr Pro Leu Ile Asp Glu Ala Asn Asp Lys Leu Ile Leu Arg Phe Gly Asp Gly Thr Ile Gln Val Arg Ser Arg Ala Asp Val Lys Asn His Ile Asp Asn Val Glu Lys Glu Met Thr Ile Asp Asn Ser Glu Asn Asp Asp Asp Arg Trp Met Gln Gly Ile Ala Val Asp Gly Asp Asp Leu Tyr Trp Leu Ser Gly Asn Ser Ser Val Asn Ser His Val Gln Ile Gly Lys Tyr Ser Leu Thr Thr Gly Gln Lys Ile Tyr Asp Tyr Pro Phe Lys Leu Ser Tyr Gln Asp Gly Ile Asn Phe Pro Arg Asp Asn Phe Lys Glu Pro Glu Gly Ile Cys Ile Tyr Thr Asn Pro Lys Thr Lys Arg Lys Ser Leu Leu Leu Ala Met Thr Asn Gly Gly Gly Gly Lys Arg Phe His Asn Leu Tyr Gly Phe Phe Gln Leu Gly Glu Tyr Glu His Phe Glu Ala Leu Arg Ala Arg Gly Ser Gln Asn Tyr Lys Leu Thr Lys Asp Asp Gly Arg Ala Leu Ser Ile Pro Asp His Ile Asp Asp Leu Asn Asp Leu Thr Gln Ala Gly Phe Tyr Tyr Ile Asp Gly Gly Thr Ala Glu Lys Leu Lys Asn Met Pro Met Asn Gly Ser Lys Arg Ile Ile Asp

Ala Gly Cys Phe Ile Asn Val Tyr Pro Thr Thr Gln Thr Leu Gly Thr

Val Gln Glu Leu Thr Arg Phe Ser Thr Gly Arg Lys Met Val Lys Met 515 520 525

Val Arg Gly Met Thr Leu Asp Val Phe Thr Leu Lys Trp Asp Tyr Gly 530 535 540

Leu Trp Thr Thr Ile Lys Thr Asp Ala Pro Tyr Gln Glu Tyr Leu Glu 545 550 555 560

Ala Ser Gln Tyr Asn Asn Trp Ile Ala Tyr Val Thr Thr Ala Gly Glu
565 570 575

Tyr Tyr Ile Thr Gly Asn Gln Met Glu Leu Phe Arg Asp Ala Pro Glu 580 585 590

Glu Ile Lys Lys Val Gly Ala Trp Leu Arg Val Ser Ser Gly Asn Ala 595 600 605

Val Gly Glu Val Arg Gln Thr Leu Glu Ala Asn Ile Ser Glu Tyr Lys 610 615 620

Glu Phe Phe Ser Asn Val Asn Ala Glu Thr Lys His Arg Glu Tyr Gly 625 630 635 640

Trp Val Ala Lys His Gln Lys 645

<210> 77

<211> 587

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 77

Met Arg Lys Leu Thr Asn Phe Lys Phe Phe Tyr Asn Thr Pro Phe Thr 1 5 10 15

Asp Tyr Gln Asn Thr Ile His Phe Asn Ser Asn Lys Glu Arg Asp Asp 20 25 30

Tyr Phe Leu Asn Gly Arg His Phe Lys Ser Leu Asp Tyr Ser Lys Gln 35 40 45

Pro Tyr Asn Phe Ile Arg Asp Arg Met Glu Ile Asn Val Asp Met Gln 50 55 60

Trp His Asp Ala Gln Gly Ile Asn Tyr Met Thr Phe Leu Ser Asp Phe 65 70 75 80

Glu Asp Arg Arg Tyr Tyr Ala Phe Val Asn Gln Ile Glu Tyr Val Asn 85 90 95

Asp Val Val Lys Ile Tyr Phe Val Ile Asp Thr Ile Met Thr Tyr Thr Gln Gly Asn Val Leu Glu Gln Leu Ser Asn Val Asn Ile Glu Arg Gln His Leu Ser Lys Arg Thr Tyr Asn Tyr Met Leu Pro Met Leu Arg Asn Asn Asp Asp Val Leu Lys Val Ser Asn Lys Asn Tyr Val Tyr Asn Gln Met Gln Gln Tyr Leu Glu Asn Leu Val Leu Phe Gln Ser Ser Ala Asp Leu Ser Lys Lys Phe Gly Thr Lys Lys Glu Pro Asn Leu Asp Thr Ser Lys Gly Thr Ile Tyr Asp Asn Ile Thr Ser Pro Val Asn Leu Tyr Val Met Glu Tyr Gly Asp Phe Ile Asn Phe Met Asp Lys Met Ser Ala Tyr Pro Trp Ile Thr Gln Asn Phe Gln Lys Val Gln Met Leu Pro Lys Asp Phe Ile Asn Thr Lys Asp Leu Glu Asp Val Lys Thr Ser Glu Lys Ile Thr Gly Leu Lys Thr Leu Lys Gln Gly Gly Lys Ser Lys Glu Trp Ser Leu Lys Asp Leu Ser Leu Ser Phe Ser Asn Leu Gln Glu Met Met Leu Ser Lys Lys Asp Glu Phe Lys His Met Ile Arg Asn Glu Tyr Met Thr Ile Glu Phe Tyr Asp Trp Asn Gly Asn Thr Met Leu Leu Asp Ala Gly Lys Ile Ser Gln Lys Thr Gly Val Lys Leu Arg Thr Lys Ser Ile Ile Gly Tyr His Asn Glu Val Arg Val Tyr Pro Val Asp Tyr Asn Ser Ala Glu Asn Asp Arg Pro Ile Leu Ala Lys Asn Lys Glu Ile Leu Ile

```
Asp Thr Gly Ser Phe Leu Asn Thr Asn Ile Thr Phe Asn Ser Phe Ala
    370
                         375
                                              380
Gln Val Pro Ile Leu Ile Asn Asn Gly Ile Leu Gly Gln Ser Gln Gln
                                                              400
                     390
385
Ala Asn Arg Gln Lys Asn Ala Glu Ser Gln Leu Ile Thr Asn Arg Ile
                                     410
Asp Asn Val Leu Asn Gly Ser Asp Pro Lys Ser Arg Phe Tyr Asp Ala
                                 425
Val Ser Val Ala Ser Asn Leu Ser Pro Thr Ala Leu Phe Gly Lys Phe
        435
                             440
Asn Glu Glu Tyr Asn Phe Tyr Lys Gln Gln Gln Ala Glu Tyr Lys Asp
    450
                         455
                                              460
Leu Ala Leu Gln Pro Pro Ser Val Thr Glu Ser Glu Met Gly Asn Ala
                     470
                                         475
                                                              480
465
Phe Gln Ile Ala Asn Ser Ile Asn Gly Leu Thr Met Lys Ile Ser Val
                 485
                                     490
Pro Ser Pro Lys Glu Ile Thr Phe Leu Gln Lys Tyr Tyr Met Leu Phe
            500
                                 505
Gly Phe Glu Val Asn Asp Tyr Asn Ser Phe Ile Glu Pro Ile Asn Ser
        515
                             520
                                                  525
Met Thr Val Cys Asn Tyr Leu Lys Cys Thr Gly Thr Tyr Thr Ile Arg
    530
                         535
                                              540
Asp Ile Asp Pro Met Leu Met Glu Gln Leu Lys Ala Ile Leu Glu Ser
545
                     550
                                         555
                                                              560
Gly Val Arg Phe Trp His Asn Asp Gly Ser Gly Asn Pro Met Leu Gln
                 565
                                     570
Asn Pro Leu Asn Asn Lys Phe Arg Glu Gly Val
            580
                                 585
<210>
       78
<211>
       487
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
·<213>
<400>
       78
Met Ile Leu Lys Arg Val Ile Thr Met Asn Asp Gln Glu Lys Ile Asp
```

Lys Phe Thr His Ser Tyr Ile Asn Asp Asp Phe Gly Leu Thr Ile Asp
20 25 30

Gln Leu Val Pro Lys Val Lys Gly Tyr Gly Arg Phe Asn Val Trp Leu 35 40 45

Gly Gly Asn Glu Ser Lys Ile Arg Gln Val Leu Lys Ala Val Lys Glu 50 60

Ile Gly Val Ser Pro Thr Leu Phe Ala Val Tyr Glu Lys Asn Glu Gly 65 70 75 80

Phe Ser Ser Gly Leu Gly Trp Leu Asn His Thr Ser Ala Arg Gly Asp 85 90 95

Tyr Leu Thr Asp Ala Lys Phe Ile Ala Arg Lys Leu Val Ser Gln Ser 100 105 110

Lys Gln Ala Gly Gln Pro Ser Trp Tyr Asp Ala Gly Asn Ile Val His
115 120 125

Phe Val Pro Gln Asp Val Gln Arg Lys Gly Asn Ala Asp Phe Ala Lys 130 135 140

Asn Met Lys Ala Gly Thr Ile Gly Arg Ala Tyr Ile Pro Leu Thr Ala 145 150 155 160

Ala Ala Thr Trp Ala Ala Tyr Tyr Pro Leu Gly Leu Lys Ala Ser Tyr 165 170 175

Asn Lys Val Gln Asn Tyr Gly Asn Pro Phe Leu Asp Gly Ala Asn Thr 180 185 190

Ile Leu Ala Trp Gly Gly Lys Leu Asp Gly Lys Gly Gly Ser Pro Ser 195 200 205

Asp Ser Ser Asp Ser Gly Ser Ser Gly Asp Ser Gly Ser Ser Leu Leu 210 215 220

Ala Leu Ala Lys Gln Ala Met Gln Glu Leu Leu Lys Lys Ile Gln Asp 225 230 235 240

Ala Leu Gln Trp Asp Val His Ser Ile Gly Ser Asp Lys Phe Phe Ser 245 250 255

Asn Asp Tyr Phe Thr Leu Glu Lys Thr Phe Asn Asn Thr Tyr His Ile 260 265 270

Lys Met Thr Ile Gly Leu Leu Asp Ser Leu Lys Lys Leu Ile Asp Ser 275 280 285

Val Gln Val Asp Ser Gly Ser Ser Ser Ser Asn Pro Thr Asp Asp Asp 290 295 300

Gly Asp His Lys Pro Ile Ser Gly Lys Ser Val Lys Pro Asn Gly Lys 305 310 315 320

Ser Gly Arg Val Ile Gly Gly Asn Trp Thr Tyr Ala Gln Leu Pro Glu 325 330 335

Lys Tyr Lys Lys Ala Ile Gly Val Pro Leu Phe Lys Lys Glu Tyr Leu 340 345 350

Tyr Lys Pro Gly Asn Ile Phe Pro Gln Thr Gly Asn Ala Gly Gln Cys 355 360 365

Thr Glu Leu Thr Trp Ala Tyr Met Ser Gln Leu His Gly Lys Arg Gln 370 375 380

Pro Thr Asp Asp Gly Gln Ile Thr Asn Gly Gln Arg Val Trp Tyr Val 385 390 395 400

Tyr Lys Lys Leu Gly Ala Lys Thr Thr His Asn Pro Thr Val Gly Tyr
405 410 415

Gly Phe Ser Ser Lys Pro Pro Tyr Leu Gln Ala Thr Ala Tyr Gly Ile 420 425 430

Gly His Thr Gly Val Val Val Ala Val Phe Glu Asp Gly Ser Phe Leu 435 440 445

Val Ala Asn Tyr Asn Val Pro Pro Tyr Val Ala Pro Ser Arg Val Val 450 455 460

Leu Tyr Thr Leu Ile Asn Gly Val Pro Asn Asn Ala Gly Asp Asn Ile 465 470 475 480

Val Phe Phe Ser Gly Ile Ala 485

<210> 79

<211> 415

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 79

Met Val Lys Gln Asn Arg Leu Asp Met Val Arg Asp Tyr Gln Asn Ala 1 5 10 15

Val Asn His Val Arg Lys Lys Ile Pro Asp Lys Tyr Asn Gln Ile Glu 20 · 25 30 Leu Val Asp Glu Leu Met Asn Asp Asp Ile Asp Tyr Tyr Ile Ser Ile 35 40 45

Ser Asn Arg Ser Asp Gly Lys Ser Phe Asn Tyr Val Ser Phe Phe Ile 50 55 60

Tyr Leu Ala Ile Lys Leu Asp Ile Lys Phe Thr Leu Leu Ser Arg His 65 70 75 80

Tyr Thr Leu Arg Asp Ala Tyr Arg Asp Phe Ile Glu Glu Ile Ile Asp 85 90 95

Glu Asn Pro Leu Phe Lys Ser Lys Arg Val Thr Phe Arg Ser Ala Arg
100 105 110

Asp Tyr Leu Ala Ile Ile Tyr Gln Asp Lys Glu Ile Gly Val Ile Thr 115 120 125

Asp Leu Asn Ser Ala Thr Asp Leu Lys Tyr His Ser Asn Phe Leu Lys 130 135 140

His Tyr Pro Ile Ile Ile Tyr Asp Glu Phe Leu Ala Leu Glu Asp Asp 145 150 155 160

Tyr Leu Ile Asp Glu Trp Asp Lys Leu Lys Thr Ile Tyr Glu Ser Ile 165 170 175

Asp Arg Asn His Gly Asn Val Asp Tyr Ile Gly Phe Pro Lys Met Phe 180 185 190

Leu Leu Gly Asn Ala Val Asn Phe Ser Ser Pro Ile Leu Ser Asn Leu 195 200 205

Asn Ile Tyr Asn Leu Leu Gln Lys His Lys Met Asn Thr Ser Arg Leu 210 215 220

Tyr Lys Asn Ile Phe Leu Glu Met Arg Arg Asn Asp Tyr Val Asn Glu 225 230 235 240

Lys Arg Asn Thr Arg Ala Phe Asn Ser Asn Asp Asp Ala Met Thr Thr 245 250 255

Gly Glu Phe Glu Phe Asn Glu Tyr Asn Leu Ala Asp Asp Asn Leu Arg 260 265 270

Asn His Ile Asn Gln Asn Gly Asp Phe Phe Tyr Ile Lys Thr Asp Asp 275 280 285

Lys Tyr Ile Lys Val Met Tyr Asn Val Thr Thr Phe Met Thr Asn Ile 290 295 300

Ile Val Val Pro Tyr Thr Lys Gln Tyr Glu Phe Cys Thr Lys Ile Arg 305 310 315 320

Asp Ile Asp Asn His Val Thr Tyr Leu Arg Asp Asp Met Phe Tyr Lys 325 330 335

Glu Asn Met Glu Arg Tyr Tyr Tyr Asn Pro Ser Asn Leu His Phe Asp 340 345 350

Asn Ala Tyr Ser Lys Asn Tyr Val Val Asp Asn Asp Arg Tyr Leu Tyr 355 360 365

Leu Asp Met Asn Lys Ile Ile Lys Phe His Ile Lys Asn Glu Met Lys 370 375 380

Lys Asn Met Ser Glu Phe Glu Arg Lys Glu Lys Ile Tyr Glu Asp Asn 385 390 395 400

Tyr Ile Glu Asn Thr Lys Lys Tyr Leu Met Lys Gln Tyr Gly Leu 405 410 415

<210> 80

<211> 408

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 80.

Met Ala Gln Gln Ser Thr Lys Asn Glu Thr Ala Leu Leu Val Ala Lys $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Ser Ala Lys Ser Ala Leu Gln Asp Phe Asn His Asp Tyr Ser Lys Ser 20 25 30

Trp Thr Phe Gly Asp Lys Trp Asp Asn Ser Asn Thr Met Phe Glu Thr 35 40 45

Phe Val Asn Lys Tyr Leu Phe Pro Lys Ile Asn Glu Thr Leu Leu Ile 50 55 60

Asp Ile Ala Leu Gly Asn Arg Phe Asn Trp Leu Ala Lys Glu Gln Asp 65 70 75 80

Phe Ile Gly Gln Tyr Ser Glu Glu Tyr Val Ile Met Asp Thr Val Pro 85 90 95

Ile Asn Met Asp Leu Ser Lys Asn Glu Glu Leu Met Leu Lys Arg Asn 100 105 110

Tyr Pro Arg Met Ala Thr Lys Leu Tyr Gly Asn Gly Ile Val Lys Lys 115 120 125

Gln Lys Phe Thr Leu Asn Asn Asn Asp Thr Arg Phe Asn Phe Gln Thr Leu Ala Asp Ala Thr Asn Tyr Ala Leu Gly Val Tyr Lys Lys Ile Ser Asp Ile Asn Val Leu Glu Glu Lys Glu Met Arg Ala Met Leu Val Asp Tyr Ser Leu Asn Gln Leu Ser Glu Thr Asn Val Arg Lys Ala Thr Ser Lys Glu Asp Leu Ala Ser Lys Val Phe Glu Ala Ile Leu Asn Leu Gln Asn Asn Ser Ala Lys Tyr Asn Glu Val His Arg Ala Ser Gly Gly Ala Ile Gly Gln Tyr Thr Thr Val Ser Lys Leu Lys Asp Ile Val Ile Leu Thr Thr Asp Ser Leu Lys Ser Tyr Leu Leu Asp Thr Lys Ile Ala Asn Thr Phe Gln Ile Ala Gly Ile Asp Phe Thr Asp His Val Ile Ser . 265 Phe Asp Asp Leu Gly Gly Val Phe Lys Val Thr Lys Glu Phe Lys Leu Gln Asn Gln Asp Ser Ile Asp Phe Leu Arg Ala Tyr Gly Asp Tyr Gln Ser Gln Leu Gly Asp Thr Ile Pro Val Gly Ala Val Phe Thr Tyr Asp Val Ser Lys Leu Lys Glu Phe Thr Gly Asn Val Glu Glu Ile Lys Pro Lys Ser Asp Leu Tyr Ala Phe Ile Leu Asp Ile Asn Ser Ile Lys Tyr Lys Arg Tyr Thr Lys Gly Met Leu Lys Pro Pro Phe His Asn Pro Glu Phe Asp Glu Val Thr His Trp Ile His Tyr Tyr Ser Phe Lys Ala Ile Ser Pro Phe Phe Asn Lys Ile Leu Ile Thr Asp Gln Asp Val Asn Pro

Lys Pro Glu Glu Glu Leu Gln Glu 405

<210> 81

<211> 327

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 81

Met Asn Asn Asp Lys Arg Gly Leu Asn Val Glu Leu Ser Lys Glu Ile 1 5 10 15

Ser Lys Arg Val Val Glu His Arg Asn Arg Phe Lys Arg Leu Met Phe 20 25 30

Asn Arg Tyr Leu Glu Phe Leu Pro Leu Leu Ile Asn Tyr Thr Asn Arg 35 40 45

Asp Thr Val Gly Ile Asp Phe Ile Gln Leu Glu Ser Ala Leu Arg Gln 50 55 60

Asn Ile Asn Val Val Gly Glu Ala Arg Asn Lys Gln Ile Met Ile 65 70 75 80

Leu Gly Tyr Val Asn Asn Thr Tyr Phe Asn Gln Ala Pro Asn Phe Ser 85 90 95

Ser Asn Phe Asn Phe Gln Phe Gln Lys Arg Leu Thr Lys Glu Asp Ile 100 105 110

Tyr Phe Ile Val Pro Asp Tyr Leu Ile Pro Asp Asp Cys Leu Gln Ile 115 120 125

His Lys Leu Tyr Asp Asn Cys Met Ser Gly Asn Phe Val Val Met Gln 130 135 140

Asn Lys Pro Ile Gln Tyr Asn Ser Asp Ile Glu Ile Ile Glu His Tyr 145 150 155 160

Thr Asp Glu Leu Ala Glu Val Ala Leu Ser Arg Phe Ser Leu Ile Met 165 170 175

Gln Ala Lys Phe Ser Lys Ile Phe Lys Ser Glu Ile Asn Asp Glu Ser 180 185 190

Ile Asn Gln Leu Val Ser Glu Ile Tyr Asn Gly Ala Pro Phe Val Lys 195 200 205

Met Ser Pro Met Phe Asn Ala Asp Asp Asp Ile Ile Asp Leu Thr Ser 210 215 220

Asn Ser Val Ile Pro Ala Leu Thr Glu Met Lys Arg Glu Tyr Gln Asn 225 230 230 240

Lys Ile Ser Glu Leu Ser Asn Tyr Leu Gly Ile Asn Ser Leu Ala Val

Asp Lys Glu Ser Gly Val Ser Asp Glu Glu Ala Lys Ser Asn Arg Gly
260 265 270

250

Phe Thr Thr Ser Asn Ser Asn Ile Tyr Leu Lys Gly Arg Glu Pro Ile 275 280 285

Thr Phe Leu Ser Lys Arg Tyr Gly Leu Asp Ile Lys Pro Tyr Tyr Asp 290 295 300

Asp Glu Thr Thr Ser Lys Ile Ser Met Val Asp Thr Leu Phe Lys Asp 305 310 315 320

Glu Ser Ser Asp Ile Asn Gly 325

245

<210> 82

<211> 251

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 82

Met Ala Arg Tyr Thr Met Thr Leu Tyr Asp Phe Ile Lys Ser Glu Leu $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ile Lys Lys Gly Phe Asn Glu Phe Val Asn Asp Asn Lys Leu Thr Phe 20 25 30

Tyr Asp Asp Glu Phe Gln Phe Met Gln Lys Met Leu Lys Phe Asp Lys 35 40 45

Asp Val Leu Ala Ile Val Asn Glu Lys Val Phe Lys Gly Phe Ser Leu 50 55 60

Lys Asp Glu Leu Ser Asp Leu Leu Phe Lys Lys Ser Phe Thr Ile His 65 70 75 80

Phe Leu Asp Arg Glu Ile Asn Arg Gln Thr Val Glu Ala Phe Gly Met 85 90 95

Gln Val Ile Thr Val Cys Ile Thr His Glu Asp Tyr Leu Asn Val Val 100 105 110

Tyr Ser Ser Ser Glu Val Glu Lys Tyr Leu Gln Ser Gln Gly Phe Thr 115 120 125

```
Glu His Asn Glu Asp Thr Thr Ser Asn Thr Asp Glu Thr Ser Asn Gln
                        135
                                             140
    130
Asn Ala Thr Ser Leu Asp Asn Ser Thr Gly Met Thr Ala Asn Arg Asn
145
                    150
                                         155
Ala Tyr Val Ser Leu Pro Gln Ser Glu Val Asn Ile Asp Val Asp Asn
                                     170
Thr Thr Leu Arg Phe Ala Asp Asn Asn Thr Ile Asp Asn Gly Lys Thr
            180
                                 185
                                                     190
Val Asn Lys Ser Ser Asn Glu Ser Asn Gln Asn Ala Lys Arg Asn Gln
        195
                             200
Asn Gln Lys Gly Asn Ala Lys Gly Thr Gln Phe Thr Lys Gln Tyr Leu
                        215
                                             220
Ile Asp Asn Ile Asp Lys Ala Tyr Asp Leu Arg Lys Lys Ile Leu Asn
                    230
                                         235
Glu Phe Asp Lys Lys Cys Phe Leu Gln Ile Trp
                245
<210>
       83
       250
<211>
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
       83
Met Lys Ser Gln Gln Gln Ala Lys Glu Trp Ile Tyr Lys His Glu Gly
                5
                                                         15
1
Ala Gly Val Asp Phe Asp Gly Ala Tyr Gly Phe Gln Cys Met Asp Leu
            20
Ser Val Ala Tyr Val Tyr Tyr Ile Thr Asp Gly Lys Val Arg Met Trp
Gly Asn Ala Lys Asp Ala Ile Asn Asn Asp Phe Lys Gly Leu Ala Thr
                        55
Val Tyr Lys Asn Thr Pro Ser Phe Lys Pro Gln Leu Gly Asp Val Ala
                    70
                                         75
                                                              80
Val Tyr Thr Asn Gly Gln Tyr Gly His Ile Gln Cys Val Leu Ser Gly
                85
Asn Leu Asp Tyr Tyr Thr Cys Leu Glu Gln Asn Trp Leu Gly Gly Gly
```

```
Phe Asp Gly Trp Glu Lys Ala Thr Ile Arg Thr His Tyr Tyr Asp Gly 115 120 125

Val Thr His Phe Ile Arg Pro Lys Phe Ser Gly Ser Asn Ser Lys Ala
```

130 135 140

Leu Glu Thr Ser Lys Val Asn Thr Phe Gly Lys Trp Lys Arg Asn Gln 145 150 155 160

Tyr Gly Thr Tyr Tyr Arg Asn Glu Asn Gly Thr Phe Thr Cys Gly Phe 165 170 175

Leu Pro Ile Phe Ala Arg Val Gly Ser Pro Lys Leu Ser Glu Pro Asn 180 185 190

Gly Tyr Trp Phe Gln Pro Asn Gly Tyr Thr Pro Tyr Asn Glu Val Cys 195 200 205

Leu Ser Asp Gly Tyr Val Trp Ile Gly Tyr Asn Trp Gln Gly Thr Arg 210 215 220

Tyr Tyr Leu Pro Val Arg Gln Trp Asn Gly Lys Thr Gly Asn Ser Tyr 225 230 235 240

Ser Val Gly Ile Pro Trp Gly Val Phe Ser 245 250

<210> 84

<211> 160

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 84

Leu Val Arg His Thr Ser Glu Met Asp Arg Trp Lys Lys Glu Arg Glu
1 5 10 15

Ala Arg Lys Glu Gln Glu Lys Asp Leu Phe Leu Asn Asp Phe Ser Asn 20 25 30

Val Asn Phe Lys Phe Asp Asp Lys Asp Leu Gln Glu Ala Tyr Ile Asp 35 40 45

Thr Trp Lys His Phe Ala His Leu Pro Tyr Phe Pro Lys Glu Arg Asn 50 55 60

Val Ser Tyr Val Asn Ala Val Ser Leu Val Arg Gly Ser Arg His Lys 65 70 75 80

Lys Leu Asn Tyr Ile Leu Glu Ile Tyr Asn Arg Asn Asp Asp Ser Asn 85 90 95

Asn Lys Asn Ala Lys Lys His Lys Tyr Ala Leu Tyr Asn Leu Gln Ala 100 105 110

Lys Asn Asn Asn Ser Ser Met Tyr Lys Tyr Ile Lys Glu Ile Asp Thr 115 120 125

Leu Tyr Lys Glu Ile Gly Lys Ser Asp Arg Pro Val Thr Asn Ile Asp 130 135 140

Asp Glu Asp Val Arg Tyr Asn Phe Leu Tyr Tyr Ala Thr Phe Asp Glu 145 150 155 160

<210> 85

<211> 122

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 85

Met Thr Asn Val Lys Asp Ile Leu Ser Arg His Gln Asn Thr Leu Ala 1 5 10 15

Arg Phe Glu Phe Glu Glu Lys Glu Arg Glu Phe Ile Lys Leu Ser Glu 20 25 30

Leu Val Glu Lys Tyr Gly Met Lys Lys Glu Tyr Ile Val Arg Ala Leu 35 40 45

Phe Thr Asn Lys Glu Ser Lys Phe Gly Glu Gln Gly Val Ile Val Thr 50 55 60

Asp Asp Tyr Asn Val Asn Leu Pro Asn His Leu Thr Glu Leu Ile Lys 70 75 80

Glu Met Arg Ala Asp Glu Asp Val Val Asp Ile Ile Asn Ala Gly Glu 85 90 95

Val Gln Phe Thr Ile Tyr Glu Tyr Glu Asn Lys Lys Gly Gln Lys Gly
100 105 110

Tyr Ser Ile Asn Phe Gly Gln Val Ser Phe 115 120

```
<210>
       86
<211>
       140
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
Met Asn Glu Val Lys Phe Arg Phe Thr Asp Ser Glu Ala Phe His Met
Phe Ile Tyr Ala Gly Asp Leu Lys Leu Leu Tyr Phe Leu Phe Val Leu
                                 25
Met Phe Val Asp Ile Ile Thr Gly Ile Ser Lys Ala Ile Lys Asn Asn
        35
Asn Leu Trp Ser Lys Lys Ser Met Arg Gly Phe Ser Lys Lys Leu Leu
                        55
Ile Phe Cys Ile Ile Ile Leu Ala Asn Ile Ile Asp Gln Ile Leu Gln
                                         75
                    70
Leu Lys Gly Gly Leu Leu Met Ile Thr Ile Phe Tyr Tyr Ile Ala Asn
                                                         95
                85
Glu Gly Leu Ser Ile Val Glu Asn Cys Ala Glu Met Asp Val Leu Val
            100
                                 105
                                                     110
Pro Glu Gln Ile Lys Asp Lys Leu Arg Val Ile Lys Asn Asp Thr Glu
        115
                            120
Lys Ser Asp Asn Asn Glu Arg Ser Arg Glu Asp Arg
                        135
<210>
       87
<211>
       136
<212>
       PRT
<213>
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
       87
Met Lys Ile Lys Thr Thr Phe Arg Leu Asn Asn Leu Ile Tyr Tyr Leu
Leu Thr Asn Arg Asp Tyr Tyr Asn Asp Lys Phe Glu Lys Phe Thr Ser
            20
                                 25
Ser Asn Lys Lys Cys Ile Val Lys Ile Asn Met Gly Asp Val Tyr Ile
        35
Glu Phe Asp Lys Gln Tyr Asp Asp Phe Glu Ile Glu Lys Glu Leu Phe
```

```
Thr Leu Asp Ile Asp Ile Asp Ile Lys Lys His Val Phe Asn Ile Leu 65 70 75 80
```

Val Phe Tyr Tyr Arg Asn Tyr Leu Ser Asn Glu Leu Ile Arg Glu Ile 85 90 95

Leu Leu Asn Val Thr Ile Asp Asp Val Leu Ser Asn Phe Asp Lys Pro 100 105 110

Leu Glu Ser Glu Leu Met Ile Ile Tyr Gln Asn Lys Val Ile Tyr Asp 115 120 125

Asn Gly Lys Val Ile Asp His Glu 130 135

<210> 88

<211> 92

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 88

Met Lys Met Val His Leu His Val Val Phe Tyr Gln Tyr Leu His Val 1 5 10 15

Ser Val Val Gln Asn Tyr Gln Asn Leu Met Ala Ile Gly Ser Asn Gln 20 25 30

Thr Val Ile His His Ile Thr Lys Phe Val Tyr Gln Met Val Thr Tyr 35 40 45

Gly Leu Val Ile Thr Gly Lys Ala His Val Ile Ile Tyr Gln Cys Ala 50 55 60

Asn Gly Met Glu Lys Gln Val Ile Val Thr Val Leu Val Phe Leu Gly 65 70 75 80

Gly Cys Ser His Asn Gly Tyr Phe Ser Leu Phe Leu 85 90

<210> 89

<211> 80

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 89

Val Thr Ile Thr Pro Cys Ser Pro Asn Phe Asp Ser Leu Phe Val Asn 1 10 15

```
Asn Ala Leu Thr Ile Tyr Ser Phe Phe Ile Pro Tyr Phe Ser Thr Asn 20 25 30
```

Ser Asp Ser Leu Ile Asn Ser Leu Ser Phe Ser Ser Asn Ser Asn Leu 35 40 45

Ala Asn Val Phe Trp Cys Leu Asp Lys Ile Ser Phe Thr Phe Val Ile 50 55 60

Leu Phe Leu Leu Leu Phe Lys Leu Phe Ala Phe Cys Asn Cys Asp Leu 65 70 75 80

<210> 90

<211> 78

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 90

Met Lys Val Asp Asp Ile Val Thr Leu Arg Val Lys Gly Tyr Ile Leu 1 5 10 15

His Tyr Leu Asp Asp Asp Asn Glu Tyr Ile Glu Glu Phe Leu Pro Leu 20 25 30

His Glu Tyr His Leu Thr Lys Thr Gln Ala Lys Glu Leu Leu Pro Asp 35 40 45

Thr Cys Lys Leu Leu Ser Thr Thr Arg Thr Thr Lys Thr Ile Gln Val 50 55 60

Tyr Tyr Asn Asp Leu Leu Gln Ile Ala Ile Ala Glu Ser Lys 65 70 75

<210> 91

<211> 73

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 91

Met Glu Arg Leu Lys Leu Leu Leu Val Tyr Arg Lys Thr Pro Leu 1 5 10 15

Ile Gln Ala Ser Ile Leu Lys Pro Leu Tyr Val Asn Asn Ser Leu Thr 20 25 30

Val Pro Leu Leu Lys Thr Ile Lys Val Ser Ile Met Ser Lys Val Gln 35 40 45

Tyr Arg Tyr Ile Arg Leu Lys Leu Lys Leu Tyr Val Val Met Tyr Met 50 55 60

```
Met Asn Ile Leu Leu Met Asn Leu Ile
                    70
<210>
       92
<211>
       70
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 92
Met Leu Ile Gly Thr Val Ser Ile Ile Thr Tyr Ser Ser Leu Tyr Cys
                                    10
Pro Ile Lys Ser Cys Ser Leu Ala Asn Gln Leu Lys Arg Leu Pro Asn
                                25
Ala Ile Ser Ile Asn Lys Val Ser Leu Ile Leu Gly Asn Lys Tyr Leu
Phe Thr Asn Val Ser Asn Ile Val Phe Glu Leu Ser His Leu Ser Pro
    50
                        55
Asn Val Gln Asp Phe Glu
<210>
       93
<211>
       68
<212>
       PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      93
Met Leu Pro Gly Leu Tyr Lys Tyr Ser Phe Leu Asn Lys Gly Thr Pro
Ile Ala Phe Leu Tyr Phe Ser Gly Asn Cys Ala Tyr Val Gln Leu Pro
                                25
Pro Ile Thr Arg Pro Leu Phe Pro Phe Gly Leu Thr Asp Leu Pro Leu
        35
                            40
                                                 45
Ile Gly Leu Trp Ser Pro Ser Ser Ser Val Gly Leu Glu Leu Leu
    50
                        55
                                            60
Pro Leu Ser Thr
65
```

```
<210>
      94
<211>
      62
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 94
Met Ser Lys Arg Phe Cys Phe Thr Met Phe Leu Leu Val Ile Val
Tyr Asp Val Val Tyr Ser Val Lys Phe Ile Arg Gln Met Leu His Asn
            20
Ile Lys Ser Tyr Thr Ser His Leu His His Gln Tyr Leu Ser Leu Val
        35
Tyr Leu Ile Tyr Gln Phe Leu Tyr Ile Lys Tyr Arg Phe Leu
                        55
<210>
      95
<211>
      60
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 95
Met His His Gln Ser Gln His Leu Pro Pro His Ala Tyr Ile Ser Ile
Leu Leu Val Val Val Ile Ser Phe Ile Ser Leu Leu Phe Leu Met
                                25
Phe Cys Tyr Pro Thr Ile Phe Thr Met Phe Cys Phe Arg Ile Asn Ile
        35
                            40
Thr Glu Glu Phe Phe Ile Phe Arg Tyr Ile Ser Leu
                       55
                                            60
    50
<210>
       96
<211>
       47
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400> 96
Met Phe Ala Lys Met Ile Ile Gln Asn Ile Asn Asn Phe Leu Glu Asn
                5
                                    10
Pro Leu Ile Asp Phe Phe Asp His Lys Leu Leu Phe Leu Ile Ala Phe
```

```
Glu Ile Pro Val Ile Ile Ser Thr Asn Ile Asn Thr Asn Lys Lys
<210>
       97
<211>
       59
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
       97
Met Arg Thr Pro Pro Lys Glu Tyr Gln His Cys Asn Tyr Tyr Leu Phe
                5
Phe His Ser Ile Gly Ala Leu Val Asn Asn Asn Val Cys Leu Ala Ser
            20
Tyr Asn Gln Ser Ile Arg Asn His Leu Ile Asn Lys Leu Arg Tyr Met
Val Tyr Asn Arg Leu Val Gly Thr Asn Ser His
                        55
<210>
       98
<211>
       58
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
Val Ser Met Tyr Ala Ser Cys Lys Ser Leu Ser Ser Asn Leu Lys Leu
Thr Leu Leu Lys Ser Phe Lys Asn Lys Ser Phe Ser Cys Ser Phe Leu
            20
                                25
                                                     30
Ala Ser Leu Ser Phe Phe His Leu Ser Ile Ser Asp Val Cys Leu Thr
        35
                            40
Asn Val Ile Asn Leu His Ile Lys His Lys
    50
                        55
<210>
       99
<211>
       58
<212>
       PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 99
Met Glu Arg Lys Tyr Lys Thr Val Leu Leu Tyr Cys Asp Glu Ile Lys
```

```
Gly His Phe Pro His Gln Ile Ser Met Phe Glu Asp Leu Tyr Asp Ala 20 25 30
```

Lys Val Val Tyr Ser Tyr Tyr Glu Tyr Asn Leu Phe Thr Lys Lys Tyr 35 40 45

Ala Tyr Ile Ile Glu Tyr Ile Lys Glu Ile 50 55

<210> 100

<211> 55

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 100

Met Asn Asn Leu Leu Asn Ile Ala Ile Val Phe Leu Leu Ala Phe Leu 1 5 10 15

Ile Thr Leu Ile Ile Leu Met Thr Leu His Ile Arg Val Ser Phe Gly 20 25 30

Val Leu Phe Thr Thr Leu Ile Ile Phe Tyr Ile Ile Phe Leu Met Val 35 40 45

Ile Tyr Ala Leu Tyr Gly Gly 50 55

<210> 101

<211> 54

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 101

Met Ile Val Tyr Ile Pro Asn Phe Ser Thr Lys Phe Ile Leu Phe Cys
1 10 15

Ile Trp Tyr Asn Asp Asn Ile Cys His Lys Ser Ser Tyr Ile Ile His 20 25 30

Asp Phe Asn Ile Phe Ile Ile Ser Phe Asp Ile Glu Glu Ile Thr Val 35 40 45

Leu Ile Asp Val Ile Ser 50

```
<210> 102
<211>
      54
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 102
Met Glu Tyr Met His Val Gln Leu Tyr Leu Leu Ser Tyr Phe Leu Gln
Asn Leu His Tyr Leu Phe Phe Val Arg Leu Val Val Gln Ser Gly Arg
            20
                                25
Cys Tyr Leu Arg His Thr Lys Thr Val Val Gln Leu Val Leu Ile Val
        35
                            40
                                                45
Ile Leu Thr Phe Leu Leu
    50
<210>
       103
<211>
      54
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 103
Val Phe Lys Trp Asn Val Asn Thr Lys Arg Tyr Tyr Ile Ala Met
Arg Leu Lys Asp Ile Phe His Ile Lys Ser Gln Cys Leu Lys Ile Tyr
Met Thr Leu Lys Leu Tyr Ile His Ile Met Asn Ile Thr Cys Ser Leu
        35
                                                45
Lys Asn Thr Arg Ile Ser
    50
<210>
      104
<211>
      53.
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400> 104
Val Asn Lys Thr Pro Asn Asp Thr Arg Ile Cys Ser Val Ile Ser Met
                                                        15
                5
                                    10
Ile Ser Val Ile Lys Asn Ala Lys Arg Lys Thr Met Ala Met Phe Asn
```

25

```
Arg Leu Phe Met Val Asn His Phe Pro Ile Ile Val Tyr Asp Phe Val
Leu Ile Asn Asn His
    50
<210>
       105
<211>
      50
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 105
Met Ile Leu Tyr Ser Ser Leu Leu Ser Ser Lys Arg Asn Lys Leu Lys
Cys Glu Arg Asn Ala Gly Met Pro Tyr Asn Pro Phe Lys Thr Thr Leu
                                25
Asp Asn Ile Thr Ser Ser Phe Glu Tyr Gly Cys Ser Leu Ile Ser Ser
                            40
Val Met
    50
<210>
      106
<211>
      49
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 106
Met Ala Cys Phe Ala Lys Ala Ser Ser Glu Leu Pro Leu Ser Pro Leu
Leu Pro Leu Ser Asp Glu Ser Leu Gly Asp Pro Pro Leu Pro Ser Asn
Leu Pro Pro Gln Ala Arg Ile Val Phe Ala Pro Ser Lys Asn Gly Leu
```

Pro

```
OWYEVESE LOSTOOT
```

```
<210> 107
<211>
      49
<212>
       PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 107
Met Pro Leu Phe Asn His Leu Tyr Gln Ile Cys Lys His Phe Leu
Ser Asn Ser Phe Lys Ile Phe Phe Leu Lys Ser Tyr Ala Leu Ser Ile
            20
                                25
Leu Ser Ile Lys Tyr Cys Leu Val Asn Cys Val Pro Phe Ala Leu Pro
        35
                                                45
Phe
<210>
      108
<211>
      48
<212>
       PRT
<213>
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
     108
Met Met Ile Leu Ile Ile Lys Thr Leu Lys Ser Ile Asn Thr Leu Tyr
Ile Ile Tyr Lys Leu Lys Ile Ile Ile Leu Gln Cys Ile Asn Ile Leu
                                25
Lys Lys Ser Ile Leu Tyr Ile Lys Lys Leu Val Asn Gln Ile Asp Gln
        35
                            40
<210>
      109
<211>
       48
<212>
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 109
Met Gln His Leu Thr Asn Lys Phe Asn Thr Val Asn Asp Ile Ile Asn
Tyr Tyr Lys Glu Gln Lys His Gly Lys Thr Lys Ser Phe Arg His Gly
                                25
Lys Arg Leu Ser Lys Cys Cys Gln Ser Cys Gln Lys Lys Asn Pro Arg
```

```
<210>
       110
<211>
       48
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<400> 110
Val Tyr Thr Ile Pro His Val Met Val Gln His Met Val Val His Tyr
                                     10
                                                         15
Ser Leu Gln Leu Lys Thr Asn His Leu Gln Lys Leu Gln Gln His
                                 25
            20
Leu Cys Asp Gln Tyr His Met Gln Leu Leu Val Ser Met Val Val Tyr
                             40
<210>
       111
<211>
       47
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
       111
Met Ser Ile Ser Asn Val Asn Asn Ser Phe Ser Ile Ser Lys Ser Ser
Tyr Cys Leu Ser Asn Ser Ile Tyr Thr Ser Pro Ile Phe Ile Phe Thr
            20
                                 25
                                                     30
Ile His Phe Leu Leu Asp Glu Val Asn Phe Ser Asn Leu Ser Leu
        35
                             40
                                                 45
<210>
       112
<211>
       47
<212>
       PRT
<213>
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
       112
Val Tyr Leu Leu His Tyr Leu Phe Asp Phe Asp Tyr Val Leu Arg Phe
                5
                                     10
Asp Tyr Phe Arg Tyr Ser Ile Tyr Ser Gln Phe Tyr Arg Tyr Gln Ser
```

Tyr Tyr Tyr Gln Arg Ile Val Thr Leu Tyr Tyr Gln His Gln Cys

```
OGVEVESE LOBIOD:
```

```
<210>
       113
<211>
       46
<212>
       PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
      113
Val Leu Tyr Leu Leu Met Met Tyr Leu Asn Leu Lys Ser Leu Leu Ala
Thr Leu Lys Lys Leu Asn Gln Asn Gln Ile Tyr Met Arg Leu Phe Trp
            20
                                25
Ile Leu Ile Gln Leu Asn Ile Asn Val Thr Gln Lys Val Cys
      114
<210>
<211>
       45
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
      114
Val Val Thr Gly His Met His Ser Tyr Gln Lys Asn Ile Lys Lys Gln
                                    10
Leu Val Tyr Leu Tyr Ser Lys Lys Asn Thr Tyr Thr Asn Gln Val Thr
Tyr Phe Leu Lys Arg Val Met Gln Asp Asn Val Gln Asn
<210>
      115
<211>
       45
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<213>
      115
<400>
Met Ser Ser Thr Phe Ile Ile Ser Leu Leu Ser Lys Lys Arg Lys
                5
                                    10
Arg Tyr Thr Phe His Lys Ile Leu Tyr Ala Tyr Ser Ile Val Leu Leu
            20
Gly His His Gln Gln Tyr Lys Thr Ile Leu Ile Leu Val
                            40
                                                 45
        35
```

```
ngymysge neino:
```

```
<210>
      116
<211>
      45
<212>
       PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 116
Met His Asp Arg Arg Leu Leu Ile Tyr Ser Phe Val Asn Leu Leu
Arg Val Met Leu Gln Ser Val His Thr His Gln Val Gly Arg Asn His
            20
Ile Asn Tyr Gly Asn Val Phe His His Arg Arg Leu Ser
        35
                            40
                                                 45
<210>
      117
<211>
      45
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      117
Met Arg Leu Val Thr Val Leu Gln His His Arg Asp Val Thr Arg Phe
                                    10
Ser Phe His His Trp Ile Asp Val Leu Ile Arg Leu Leu Tyr His Asp
                                25
His Pro Val Gln Tyr Ala Cys Leu Lys Leu Ser His His
                            40
<210>
       118
<211>
       44
<212>
       PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      118
Met Leu Pro Ile Tyr Val Met Ile Cys Phe Ile Lys Lys Thr Trp Asn
                5
                                    10
                                                         15
Val Ile Thr Thr Ile Gln Ala Ile Tyr Ile Leu Thr Met Leu Thr Leu
            20
                                25
                                                     30
Lys Ile Thr Trp Leu Ile Met Ile Asp Ile Tyr Ile
        35
                            40
```

```
DG7E789E . D81001
```

```
<210> 119
<211>
      42
<212>
      PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 119
Met Ala Pro Ser Lys Asn Cys Ser Arg Thr Lys Val Ser Lys Ser Thr
Leu Val Ser Lys Ala Phe Phe Gly Ile Pro Ala Glu Ala Ile Leu Ile
            20
                                25
Phe Pro Phe Thr Ser Tyr Ala Tyr Phe Leu
        35
<210>
      120
<211>
      42
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      120
Met Ile Ile His Leu Ser Tyr His Ile Lys Thr Val Leu Ile Ser His
                5
                                    10
Val Ile Thr Leu Lys Ser Leu Arg Val Phe Ala Phe Ile Gln Ile Gln
                                25
Lys Gln Asn Val Asn Arg Tyr Tyr Leu Leu
        35
                            40
<210>
      121
<211>
      42
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      121
Met Asn Val Cys Lys Leu Phe Arg Cys Glu Phe Cys Lys Thr Phe His
                5
                                    10
Ser Ile Val Ile Gly Phe Thr Ile Ile His Ile Ile Ile Phe Ile Lys
            20
                                25
                                                     30
Asn Arg Ile Ile Lys Ile Cys Phe Lys Leu
        35
                            40
```

```
COLEGO, HOSTACOS
```

```
<210>
      122
<211>
       42
<212>
       PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400>
      122
Met Arg Thr Leu Leu Thr Leu Ser Met Leu Glu Lys Phe Asn Ser Gln
Phe Met Asn Met Lys Thr Lys Lys Val Lys Lys Val Thr Gln Ser Ile
                                 25
            20
Leu Val Lys Tyr His Phe Asn Thr Ile Ser.
<210>
       123
<211>
       41
<212>
       PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
       123
Met Arg Gly Gln Val Leu Thr Leu Met Val His Met Asp Phe Asn Val
                                     10
Trp Thr Tyr Gln Leu Leu Met Cys Ile Thr Leu Leu Thr Val Lys Phe
                                 25
Ala Cys Gly Val Met Leu Lys Thr Arg
<210>
       124
<211>
       41
<212>
       PRT
<213>
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      124
Val Cys Tyr Val Phe His Ser Arg Asn Arg Phe Val Ala Phe Leu Lys
                5
                                     10
Lys Cys Phe Cys Lys Val Leu Met Tyr Ser Phe Tyr Ala Phe Val Ile
                                 25
            20
Asn Cys Ile Tyr Leu Asn Trp Ile Ile
        35
```

```
<210>
      125
<211>
      40
<212>
       PRT
<213>
      Staphylococcus aureus Bacteriophage 44 AHJD
<400> 125
Met Ile Thr Met Asn Tyr Thr Ile Ser Leu Thr Val Thr Lys Thr Leu
Asn Val Ile Tyr Tyr Ser Leu His Leu Ser His His Val His Cys Ile
            20
                                25
Thr Tyr Trp Phe Leu Ser Asn Thr
        35
<210> 126
<211>
      40
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      126
Met Ile Leu Val Met Leu Ile Leu Asn Leu Met Ile Lys Ile Tyr Lys
                5
                                    10
Arg Arg Thr Leu Thr His Gly Asn Ile Leu His Ile Cys Pro Ile Phe
                                25
Leu Lys Lys Glu Thr Tyr His Met
        35
<210>
      127
<211>
      39
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      127
Met Trp Phe Ile His Gln Val Lys Leu Lys Asn Thr Tyr Asn His Lys
                5
                                    10
Ala Ser Gln Asn Thr Met Lys Ile Gln Gln Val Thr Leu Met Lys His
            20
                                                     30
                                25
Arg Ile Lys Met Leu His Leu
        35
```

```
<210>
          128
    <211>
           3.9
    <212>
          PRT
    <213>
          Staphylococcus aureus Bacteriophage 44 AHJD
    <400> 128
    Met Thr Gly Met Glu Ile Arg Cys Tyr Ser Thr Leu Val Arg Phe His
    Lys Lys Leu Val Leu Ser Tyr Val Gln Asn Gln Leu Leu Val Ile Ile
                20
    Met Lys Phe Glu Tyr Ile Gln
            35
    <210> 129
    <211>
           38
    <212>
          PRT
    <213> Staphylococcus aureus Bacteriophage 44 AHJD
    <400> 129
IJ
    Met Cys Leu Val Ile Ile Leu Leu Leu Val Phe Trp Leu Asn Asp Thr
17. H
П
    Arg Glu Val Val Lys Ile Pro Gln Cys Ile His Tyr His His Leu Ser
IJ
                                     25
Asn Glu Val Tyr Asn Leu
O
            35
    <210>
          130
    <211>
           37
    <212>
           PRT
           Staphylococcus aureus Bacteriophage 44 AHJD
    <400>
          130
    Val Ser Ile Thr Leu Gln Val Thr Lys Trp Asn Tyr Leu Glu Thr Arg
                                         10
    Gln Lys Lys Leu Lys Lys Trp Val His Gly Tyr Val Cys Gln Val Val
                20
                                                         30
                                     25
    Thr Gln Ser Val Lys
            35
```

```
. OSTATE DELDI
```

```
<210>
      131
<211>
       36
<212>
       PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 131
Met Tyr His His Met Leu His His His Val Trp Tyr Cys Ile His Ser
Leu Met Ala Tyr Gln Ile Met Leu Val Ile Ile Leu Tyr Ser Leu Val
            20
Val Leu Leu Asn
        35
<210>
       132
<211>
       36
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      132
Met His Ile Ser Tyr Asp Ser Val Gln Thr Ser Tyr Leu Ser Val Arg
Phe Gln Tyr Pro Ile Tyr Leu Arg Leu Ser Gly Arg Ile Asn Trp Gly
                                25
Ser Ile Arg Val
        35
<210>
      133
<211>
       36
<212>
       PRT
<213>
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      133
Met Asp Phe Val Thr Leu Asp Tyr Leu Asn Arg His Tyr Ala Lys Ile
                                                         15
Leu His Gln Ile Leu Lys Leu Leu Leu Ile Val Pro Leu Thr Trp Gly
            20
Arg Cys His Val
        35
```

```
OSTETE DEIDL
```

```
<210> 134
<211>
      36
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 134
Met Tyr His Phe His Phe Tyr Asn Met Cys Arg Ile Gly Phe Val Ser
Ile Phe Gln Met Tyr Leu Leu Met Phe Leu Met Leu Cys Tyr Tyr
Tyr Leu Lys Ile
        35
      135
<210>
<211>
      35
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400> 135
Met Cys Phe Gly Val Leu Ile Lys Tyr Leu Leu Arg Leu Ser Phe Tyr
                5
                                    10 -
Phe Ser Ser Tyr Leu Asn Tyr Leu Leu Ser Ala Ile Ala Ile Cys Ser
                                25
Lys Ser Leu
        35
<210>
      136
<211>
      34
<212> PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      136
Val Val Phe Ala Thr Gln Leu Thr Asn Leu Leu Ile Leu Ile Lys Lys
                5
                                    10
                                                        15
Gln Ile Thr Cys Thr Leu His Asn Pro Ile Leu Lys Asn Leu Lys Val
            20
                                25
Phe Gly
```

```
<210>
      137
<211>
      34
<212>
      PRT
      Staphylococcus aureus Bacteriophage 44 AHJD
<213>
<400> 137
Met Arg Leu Val Phe Phe Leu Ile Ile Leu Ala Trp Leu Val Leu Leu
Lys Arg Val Val Asn Tyr His Cys His His Tyr Tyr His Cys Gln Thr
                                                    30
Asn His
<210>
       138
<211>
       33
<212>
      PRT
<213> Staphylococcus aureus Bacteriophage 44 AHJD
<400> 138
Met Thr Ser Gln Ser Ile Asn Leu Cys Pro Lys Tyr Ile Thr Val His
His Leu Leu Lys Cys His Leu Cys Leu Met Gln Met Thr Ile Ser Leu
                                25
Ile
<210>
      139
<211>
      33
<212>
       PRT
       Staphylococcus aureus Bacteriophage 44 AHJD
<400>
      139
Val Val Glu Asn Val Ser Ile Ile Tyr Met Val Ser Ser Asn Leu Val
                                    10
Ser Met Asn Thr Leu Lys His Tyr Ala Gln Glu Val His Lys Thr Ile
            20
```

Asn

<210> 140

<211> 33

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 140

Met Ile Phe Phe Ile Leu Lys Val Thr Ser Val His Phe His Leu Thr 1 5 10 15

Thr Tyr Phe Gln Leu Asn Val Gln Tyr Ile Thr Asn Leu Ile Cys Ile 20 25 30

Tyr

<210> 141

<211> 133

<212> PRT

<213> Staphylococcus aureus Bacteriophage 44 AHJD

<400> 141

Met Thr Glu Phe Asp Glu Ile Val Lys Pro Asp Asp Lys Glu Glu Thr $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Glu Ser Thr Glu Glu Asn Leu Glu Ser Thr Glu Glu Thr Ser Glu 20 25 30

Ser Thr Glu Glu Ser Thr Glu Glu Ser Thr Glu Glu Ser Thr Glu Asp 35 40 45

Lys Thr Val Glu Thr Ile Glu Glu Glu Asn Glu Asn Lys Leu Glu Pro 50 55 60

Thr Thr Thr Asp Glu Asp Ser Ser Lys Phe Asp Pro Val Val Leu Glu 65 70 . 75 80

Gln Arg Ile Ala Ser Leu Glu Gln Gln Val Thr Thr Phe Leu Ser Ser 85 90 95

Gln Met Gln Gln Pro Gln Gln Val Gln Gln Thr Gln Ser Asp Val Thr 100 105 110

Glu Ser Asn Lys Glu Asp Asn Asp Tyr Ser Asp Glu Glu Leu Val Asp 115 120 125

Lys Leu Asp Leu Asp 130

<210> 142 <211> 100 <212> PRT Staphylococcus aureus Bacteriophage 44 AHJD <213> <400> 142 Met Val Asn Val Asp Asn Ala Pro Glu Glu Lys Gly Gln Ala Tyr Thr Glu Met Leu Gln Leu Phe Asn Lys Leu Ile Gln Trp Asn Pro Ala Tyr 20 25 Thr Phe Asp Asn Ala Ile Asn Leu Leu Ser Ala Cys Gln Gln Leu Leu 35 45 Leu Asn Tyr Asn Ser Ser Val Val Gln Phe Leu Asn Asp Glu Leu Asn 55 Asn Glu Thr Lys Pro Glu Ser Ile Leu Ser Tyr Ile Ala Gly Asp Asp Pro Ile Glu Gln Trp Asn Met His Lys Gly Phe Tyr Glu Thr Tyr Asn 85 Val Tyr Val Phe 100 <210> 143 <211> 65 <212> PRT <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 143 Met Glu Asn Glu Thr Lys Asn Ile Glu Leu Lys His Val Phe Arg Phe 5 15 Lys Asn Gly Ser Leu Cys Ile Ala Leu Phe Asp Arg Thr Glu Asn Glu 25 Ile Ser Phe Tyr Asp Val Asp Ile Asp Glu Ile Glu Asp Leu Asn His Asn Ser Val Leu Arg Val Ile Ser Thr Leu Leu Gly Ser Asp Asn Asn Gly

<210> 144 <211> 60 <212> PRT <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 144 Met Tyr Glu Gly Asn Asn Met Arg Ser Met Met Gly Thr Ser Tyr Glu Asp Ser Arg Leu Asn Lys Arg Thr Glu Leu Asn Glu Asn Met Ser Ile 20 25 Asp Thr Asn Lys Ser Glu Asp Ser Tyr Gly Val Gln Ile His Ser Leu 35 Ser Lys Gln Ser Phe Thr Gly Asp Val Glu Glu Glu 55 <210> 145 <211> 49 <212> PRT <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 145 Met Lys Thr Cys Gln Leu Ile Gln Ile Lys Val Lys Ile Val Met Val Tyr Lys Phe Ile His Phe Gln Asn Asn His Leu Gln Val Thr Leu Arg 25 Arg Asn Asn Lys Leu Trp His Asn Asn Leu Gln Lys Met Lys Leu His 35 Phe <210> 146 <211> 43 <212> PRT <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 146

79

10

30

Met Ile Val Leu Lys Val Asn Glu Phe Val His His Asn Tyr Leu His

Phe Tyr Leu Tyr Gln Leu Thr Cys Phe His Leu Ile Leu Phe Val Tyr

25

Leu Ile	Leu 35	Asn	Leu	His	Met	Met 40	Tyr	Pro	Ser						
<211> 3 <212> F	147 36 PRT Staph	wl oc	.0.001	16 31	120116	· Pa	, atori	onh	age /	1 <i>1</i> 7 1	חד ג				
<213/ S	ocapi.	IŽIOC	occi	is at	ireus	р Ба	cceri	горпа	ige -	A PE	עטו				
<400> 1	L47														
Met Phe	Ser	Phe	Asn 5	Ser	Val	Arg	Leu	Phe 10	Asn	Leu	Glu	Ser	Ser 15	Tyr	
Asp Val	Pro	Ile 20	Ile	Glu	Arg	Met	Leu 25	Phe	Pro	Ser	Tyr	Met 30	Phe	Lys	
Phe Leu	Leu 35	Ile											٠.		
<211> 5 <212> 0	148 53 DNA Staph	nyloc	coccu	ıs aı	ıreus	s Bac	cteri	opha	ige 4	14 AI	HJD				
<400> 1 gatcccgg	l48 gtc g	jacca	agct	it ta	accca	itac	g aco	jtcco	caga	cta	cgcca	agc 1	tga		53
<211> 5 <212> 0	149 53 DNA Staph	nyloc	occu	ıs aı	ıreus	s Bac	cteri	opha	ige 4	14 AI	HJD				
<400> 1	l49 gct g	ıgcgt	agto	t g	ggaco	gtcgi	t ato	ggta	aaag	ctt	ggtc	gac (cgg		53
<211> 2 <212> D	150 21 ONA Staph	nyloc	occu	ıs aı	ıreus	s Bac	cteri	opha	age 4	14 AI	HJD				
<400> 1	150 gag t	aaaa	ıtaac	a t											21

<211>	37		
	31		
<212>	DNA		
<213>	Staphylococcus a	aureus Bacteriophage 44 AHJD	
<400>	151		
cgggat	ccgc ctccttttct (caacagtcac ctgattt	37
2010 >	1 5 0	•	
		·	
		aureus Bacteriophage 44 AHJD	
\213/	Scapily Lococcus &	aureus bacterrophage 44 Anob	
<400>	152		
cgggat	ccat gaggggttcc (gaagacg	27
4010 5	1.50		
		aurous Bastoriophago 44 AUID	
\213/	Staphyrococcus a	aureus bacterrophage 44 Anob	
<400>	153		
		tttc	24
_	33		
	•		
<210>	154		
<211>	27		
	DNA		
<213>	Staphylococcus a	aureus Bacteriophage 44 AHJD	
ccgctc	gage tecaaattee a	aaaacag	27
<210>	155		
<211>	26		
<212>	DNA		
<213>	Staphylococcus a	aureus Bacteriophage 44 AHJD	
<400>	155		
		tttac	26
-555000			_ `
	<213> <400> cgggatc <210> <211> <212> <213> <400> cgggatc <210> <211> <212> <213> <400> cccaagc <210> <211> <212> <213> <400> cccaagc <210> <211> <212> <213> <400> ccgctcg <210> <211> <212> <213> <400> ccgctcg <210> <211> <212> <213> <400> ccgctcg <210> <210> <211> <212> <213> <400> ccgctcg <210> <210> <210> <210> <211> <212> <213> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210> <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <210 <2	<pre><213> Staphylococcus <400> 151 cgggatccgc ctcctttct <210> 152 <211> 27 <212> DNA <213> Staphylococcus <400> 152 cgggatccat gaggggttcc <210> 153 <211> 24 <212> DNA <213> Staphylococcus <400> 153 cccaagctta caatttggac <400> 153 cccaagctta caatttggac <210> 154 <211> 27 <212> DNA <213> Staphylococcus <400> 154 ccgctcgagc tccaaattcc <210> 154 ccgctcgagc tccaaattcc <210> 155 <211> 26 <212> DNA <213> Staphylococcus <400> 155 <211> 26 <210> 155 <211> 26 <212> DNA <213> Staphylococcus <400> 155</pre>	<pre><213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 151 cgggatccgc ctccttttct caacagtcac ctgattt <210> 152 <211> 27 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 152 cgggatccat gaggggttcc gaagacg <210> 153 <211> 24 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 153 cccaagctta caatttggac tttc <210> 154 <211> 27 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 154 cccaagctta caatttggac tttc <210> 154 cccaagctta caattcc aaaacag <210> 155 <211> 26 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD <400> 154 ccgctcgagc tccaaattcc aaaacag</pre> <210> 155 <211> 26 <212> DNA <213> Staphylococcus aureus Bacteriophage 44 AHJD <513> Staphylococcus aureus Bacteriophage 44 AHJD <514

<210		156														
<211		10														
<212 <213		DNA Stap	hylo	cocci	us a	ureu	s Ba	cter	ioph	age	44 A	HJD		,	-	
<400	15	156														
gcgt																10
gcgt	Lega	ccg				•										10
<210)>	157														
<211	L> :	20														
<212	2>	DNA														
<213	3>	Stap	hylo	cocci	us a	ureu	s Ba	cter	ioph	age	44 A	HJD				
<400)>	157														
tatt	atc	caa	aact	tgaad	ca											20
<210		158									-					
<211		20														
<212		DNA				•										
<213	3> ;	Stap	hylo	coccı	ıs aı	ureus	s Ba	cter	ioph	age '	44 A	HJD				
<400		158														
cggt	ggta	ata 1	tcca	gtgat	:t											20
<210)> :	159														
<211		133														
<212		PRT					_									
<213	> :	stapi	путо	cocci	ıs aı	ureus	в Ва	cter	lopha	age 4	44 A)	HJD				
<400	> :	159														
Met	Lys	Ile	Lys	Val	Lys	Lys	Glu	Met	Arg	Leu	Asp	Glu	Leu	Ile	Lys	
L .				5					10	•				15		
rp.	Ala	Arg	Glu	Asn	Pro	Asp	Leu	Ser	Gln	Cys	Lys	Ile	Phe	Phe	Ser	
			20					25					30			
hr	Gly	Phe	Ser	Asp	Gly	Phe	Val	Arg	Phe	His	Pro	Asn	Thr	Asn	Lys	
		35					40					45				
Cys	Ser	Thr	Ser	Ser	Phe	Ile	Pro	Ile	Asp	Ile	Pro	Phe	Ile	Val	Asp	
	50					55			•		60				•	
:le	Gl 11	Lvs	Glu	Val	Thr	Glu	Gl 11	Thr	Lvs	Val	Asp	Ara	Len	Tle	Glu	
65	J_ u	_ _J	J= U		70	<u></u>	JIU		ر بر	75		9	u		80	
-					. •				•							
Leu	Phe	Glu	Ile	Gln	Glu	Gly	Asp	Tyr		Ser	Thr	Leu	Tyr		Asn	
				25					വ					95		

Thr Ser Ile Lys Glu Cys Leu Tyr Gly Arg Cys Val Pro Thr Lys Ala 100 105 110



Phe Tyr Ile Leu Asn Asp Asp Leu Thr Met Thr Leu Ile Trp Lys Asp 115 120 125

Gly Glu Leu Leu Val 130

BEST AVAILABLE COPY